

GenCore version 5.1.6
Copyright (c) 1993 - 2004 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: March 5, 2004, 16:09:43 ; Search time 33 Seconds
(without alignments)
19.196 Million cell updates/sec

Title: US-09-998-491-9
Perfect score: 15
Sequence: 1 RER 3

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 809742 seqs, 211153259 residues

Total number of hits satisfying chosen parameters: 441

Minimum DB seq length: 0
Maximum DB seq length: 20

Post-processing: Minimum Match 100%
Maximum Match 100%
Listing first 1000 summaries

Database : Published Applications AA.*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep.*
2: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep.*
3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep.*
4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep.*
5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep.*
6: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep.*
7: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep.*
8: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep.*
9: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep.*
10: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep.*
11: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep.*
12: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep.*
13: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep.*
14: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep.*
15: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep.*
16: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep.*
17: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep.*
18: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES			
Result No.	Score	Query Match Length DB ID	Description
1	15	100.0	3 10 US-09-998-491-9 Sequence 9, Appl
2	15	100.0	4 10 US-09-998-491-10 Sequence 10, Appl
3	15	100.0	4 10 US-09-998-491-11 Sequence 11, Appl
4	15	100.0	4 14 US-10-357-467-29 Sequence 29, Appl
5	15	100.0	5 9 US-09-096-749A-59 Sequence 59, Appl
6	15	100.0	5 10 US-09-903-412-59 Sequence 59, Appl
7	15	100.0	5 10 US-09-998-491-3 Sequence 3, Appl
8	15	100.0	5 10 US-09-998-491-4 Sequence 4, Appl
9	15	100.0	5 14 US-10-174-717A-59 Sequence 59, Appl
10	15	100.0	5 14 US-10-165-155-59 Sequence 59, Appl
11	15	100.0	5 14 US-10-190-162-59 Sequence 59, Appl
12	15	100.0	6 14 US-10-082-747A-60 Sequence 60, Appl
13	15	100.0	6 14 US-10-315-964A-462 Sequence 462, Appl
14	15	100.0	6 14 US-10-315-964A-524 Sequence 524, Appl
15	15	100.0	6 14 US-10-315-964A-525 Sequence 525, Appl

Sequence 462, App
Sequence 524, App
Sequence 525, App
Sequence 462, App
Sequence 524, App
Sequence 525, App
Sequence 21, Appl
Sequence 2, Appl
Sequence 495, App
Sequence 508, App
Sequence 17, Appl
Sequence 49, Appl
Sequence 45, App
Sequence 503, App
Sequence 506, App
Sequence 519, App
Sequence 39, Appl
Sequence 64, Appl
Sequence 91, Appl
Sequence 111, Appl
Sequence 54, Appl
Sequence 54, Appl
Sequence 2180, App
Sequence 2181, App
Sequence 39, Appl
Sequence 64, Appl
Sequence 91, Appl
Sequence 111, Appl
Sequence 79, Appl
Sequence 80, Appl
Sequence 516, App
Sequence 621, App
Sequence 379, App
Sequence 1036, App
Sequence 41, Appl
Sequence 68, Appl
Sequence 75, Appl
Sequence 94, Appl
Sequence 112, App
Sequence 103, App
Sequence 187, App
Sequence 5, Appl
Sequence 15, Appl
Sequence 2185, App
Sequence 2186, App
Sequence 2187, App
Sequence 6, Appl
Sequence 429, App
Sequence 31, Appl
Sequence 41, Appl
Sequence 68, Appl
Sequence 75, Appl
Sequence 94, Appl
Sequence 112, App
Sequence 127, App
Sequence 127, App
Sequence 150, App
Sequence 139, App
Sequence 696, App
Sequence 1224, App
Sequence 1231, App
Sequence 1264, App
Sequence 1734, App
Sequence 1742, App
Sequence 1746, App
Sequence 1789, App
Sequence 2342, App
Sequence 2818, App
Sequence 2836, App
Sequence 2864, App
Sequence 3369, App
Sequence 3427, App
Sequence 3432, App

89	15	100.0	9	15	US-10-107-532-4399	Sequence 4399, Ap	162	10	15	US-10-107-532-3096	Sequence 3096, Ap
90	15	100.0	9	15	US-10-107-532-4451	Sequence 4451, Ap	163	10	15	US-10-107-532-3115	Sequence 3115, Ap
91	15	100.0	9	15	US-10-107-532-4469	Sequence 4469, Ap	164	10	15	US-10-107-532-3131	Sequence 3131, Ap
92	15	100.0	9	15	US-10-107-532-4530	Sequence 4530, Ap	165	10	15	US-10-107-532-3138	Sequence 3138, Ap
93	15	100.0	9	15	US-10-107-532-4568	Sequence 4568, Ap	166	10	15	US-10-107-532-3139	Sequence 3139, Ap
94	15	100.0	9	15	US-10-107-532-4573	Sequence 4573, Ap	167	10	15	US-10-107-532-3153	Sequence 3153, Ap
95	15	100.0	9	15	US-10-107-532-4587	Sequence 4587, Ap	168	10	15	US-10-107-532-3163	Sequence 3163, Ap
96	15	100.0	9	15	US-10-107-532-4586	Sequence 4586, Ap	169	10	15	US-10-107-532-3614	Sequence 3614, Ap
97	15	100.0	9	15	US-10-107-532-4627	Sequence 4627, Ap	170	10	15	US-10-107-532-3629	Sequence 3629, Ap
98	15	100.0	9	15	US-10-107-532-4642	Sequence 4642, Ap	171	10	15	US-10-107-532-3689	Sequence 3689, Ap
99	15	100.0	10	9	US-09-731-221-7	Sequence 7, Appl	172	10	15	US-10-107-532-3311	Sequence 3311, Ap
100	15	100.0	10	10	US-09-983-802-388	Sequence 388, App	173	10	15	US-10-107-532-5398	Sequence 5398, Ap
101	15	100.0	10	10	US-09-894-912A-21	Sequence 21, Appl	174	10	15	US-10-107-532-5404	Sequence 5404, Ap
102	15	100.0	10	10	US-09-809-638-208	Sequence 208, App	175	10	15	US-10-107-532-5450	Sequence 5450, Ap
103	15	100.0	10	10	US-09-809-638-394	Sequence 394, App	176	10	15	US-10-452-510-65	Sequence 65, Appl
104	15	100.0	10	10	US-09-809-638-471	Sequence 471, App	177	10	15	US-09-966-871-3	Sequence 3, Appl
105	15	100.0	10	10	US-09-809-638-566	Sequence 566, App	178	10	15	US-09-876-904A-239	Sequence 239, App
106	15	100.0	10	10	US-09-809-638-671	Sequence 671, App	179	10	15	US-09-940-316B-57	Sequence 57, Appl
107	15	100.0	10	10	US-09-876-904A-231	Sequence 231, App	180	10	15	US-10-039-645-3	Sequence 3, Appl
108	15	100.0	10	10	US-09-876-904A-238	Sequence 238, App	181	10	15	US-10-062-710-165	Sequence 165, App
109	15	100.0	10	10	US-09-572-404B-214	Sequence 214, App	182	10	15	US-10-062-710-166	Sequence 166, App
110	15	100.0	10	10	US-09-572-404B-216	Sequence 216, App	183	10	15	US-10-139-084-3	Sequence 3, Appl
111	15	100.0	10	10	US-09-572-404B-1630	Sequence 1630, App	184	10	15	US-10-224-999A-2198	Sequence 2198, Ap
112	15	100.0	10	10	US-09-572-404B-1801	Sequence 1801, Ap	185	10	15	US-10-224-999A-2199	Sequence 2199, Ap
113	15	100.0	10	10	US-09-572-404B-1809	Sequence 1809, Ap	186	10	15	US-10-224-999A-2200	Sequence 2200, Ap
114	15	100.0	10	10	US-09-572-404B-1811	Sequence 1811, Ap	187	10	15	US-10-224-999A-2201	Sequence 2201, Ap
115	15	100.0	10	10	US-09-572-404B-1823	Sequence 1823, Ap	188	10	15	US-10-224-999A-2202	Sequence 2202, Ap
116	15	100.0	10	10	US-09-572-404B-1825	Sequence 1825, Ap	189	10	15	US-10-411-869A-30	Sequence 30, Appl
117	15	100.0	10	10	US-09-572-404B-2048	Sequence 2048, Ap	190	10	15	US-08-666-340-4	Sequence 4, Appl
118	15	100.0	10	10	US-09-572-404B-3142	Sequence 3142, Ap	191	10	15	US-09-096-749A-118	Sequence 118, App
119	15	100.0	10	10	US-09-572-270A-26	Sequence 26, Appl	192	10	15	US-09-903-412-118	Sequence 118, App
120	15	100.0	10	10	US-09-572-270A-120	Sequence 120, App	193	10	15	US-10-174-717A-118	Sequence 118, App
121	15	100.0	10	10	US-09-572-270A-467	Sequence 467, App	194	10	15	US-10-165-155-118	Sequence 118, App
122	15	100.0	10	10	US-09-572-270A-654	Sequence 654, App	195	10	15	US-10-286-457-204	Sequence 204, App
123	15	100.0	10	11	US-09-933-780C-36	Sequence 36, Appl	196	10	15	US-10-190-162-118	Sequence 118, App
124	15	100.0	10	14	US-10-228-806-13	Sequence 13, Appl	197	10	15	US-10-224-999A-2206	Sequence 2206, Ap
125	15	100.0	10	14	US-10-228-806-61	Sequence 61, Appl	198	10	15	US-10-224-999A-2207	Sequence 2207, Ap
126	15	100.0	10	14	US-10-228-806-62	Sequence 62, Appl	199	10	15	US-10-224-999A-2208	Sequence 2208, Ap
127	15	100.0	10	14	US-10-228-806-63	Sequence 63, Appl	200	10	15	US-10-224-999A-2209	Sequence 2209, Ap
128	15	100.0	10	14	US-10-228-806-64	Sequence 64, Appl	201	10	15	US-10-224-999A-2210	Sequence 2210, Ap
129	15	100.0	10	14	US-10-228-806-65	Sequence 65, Appl	202	10	15	US-10-224-999A-2211	Sequence 2211, Ap
130	15	100.0	10	14	US-10-190-082-339	Sequence 339, App	203	10	15	US-10-452-510-63	Sequence 63, Appl
131	15	100.0	10	14	US-10-167-831-32	Sequence 32, Appl	204	10	15	US-10-452-510-64	Sequence 64, Appl
132	15	100.0	10	14	US-10-167-831-44	Sequence 44, Appl	205	10	15	US-09-999-220B-21	Sequence 21, Appl
133	15	100.0	10	14	US-10-224-999A-2191	Sequence 2191, Ap	206	10	15	US-09-999-220B-47	Sequence 47, Appl
134	15	100.0	10	14	US-10-224-999A-2192	Sequence 2192, Ap	207	10	15	US-09-999-220B-73	Sequence 73, Appl
135	15	100.0	10	14	US-10-224-999A-2193	Sequence 2193, Ap	208	10	15	US-10-014-162-63	Sequence 63, Appl
136	15	100.0	10	14	US-10-224-999A-2194	Sequence 2194, Ap	209	10	15	US-10-120-604-36	Sequence 36, Appl
137	15	100.0	10	14	US-10-155-693-28	Sequence 28, Appl	210	10	15	US-10-120-604-64	Sequence 64, Appl
138	15	100.0	10	14	US-10-200-708-104	Sequence 104, App	211	10	15	US-10-224-999A-2215	Sequence 2215, Ap
139	15	100.0	10	15	US-10-200-708-260	Sequence 260, App	212	10	15	US-10-224-999A-2216	Sequence 2216, Ap
140	15	100.0	10	15	US-10-100-303A-15	Sequence 15, Appl	213	10	15	US-10-224-999A-2217	Sequence 2217, Ap
141	15	100.0	10	15	US-10-100-303A-48	Sequence 48, Appl	214	10	15	US-10-224-999A-2218	Sequence 2218, Ap
142	15	100.0	10	15	US-10-100-303A-49	Sequence 49, Appl	215	10	15	US-10-224-999A-2219	Sequence 2219, Ap
143	15	100.0	10	15	US-10-100-303A-50	Sequence 50, Appl	216	10	15	US-10-224-999A-2220	Sequence 2220, Ap
144	15	100.0	10	15	US-10-100-303A-51	Sequence 51, Appl	217	10	15	US-10-224-999A-2221	Sequence 2221, Ap
145	15	100.0	10	15	US-10-100-303A-52	Sequence 52, Appl	218	10	15	US-10-224-999A-2222	Sequence 60, Appl
146	15	100.0	10	15	US-10-107-532-422	Sequence 422, App	219	10	15	US-10-193-477-60	Sequence 60, Appl
147	15	100.0	10	15	US-10-107-532-434	Sequence 434, App	220	10	15	US-10-193-477-61	Sequence 1, Appl
148	15	100.0	10	15	US-10-107-532-456	Sequence 456, App	221	10	15	US-09-096-749A-1	Sequence 1, Appl
149	15	100.0	10	15	US-10-107-532-461	Sequence 461, App	222	10	15	US-09-876-187-23	Sequence 23, Appl
150	15	100.0	10	15	US-10-107-532-480	Sequence 480, App	223	10	15	US-09-900-530A-19	Sequence 19, Appl
151	15	100.0	10	15	US-10-107-532-966	Sequence 966, App	224	10	15	US-09-903-412-1	Sequence 1, Appl
152	15	100.0	10	15	US-10-107-532-1480	Sequence 1480, Ap	225	10	15	US-09-992-665-3	Sequence 3, Appl
153	15	100.0	10	15	US-10-107-532-1490	Sequence 1490, Ap	226	10	15	US-10-078-547-27	Sequence 27, Appl
154	15	100.0	10	15	US-10-107-532-1524	Sequence 1524, Ap	227	10	15	US-10-014-162-62	Sequence 62, Appl
155	15	100.0	10	15	US-10-107-532-1533	Sequence 1533, Ap	228	10	15	US-10-012-456A-42	Sequence 42, Appl
156	15	100.0	10	15	US-10-107-532-1552	Sequence 1552, Ap	229	10	15	US-10-174-717A-1	Sequence 1, Appl
157	15	100.0	10	15	US-10-107-532-1999	Sequence 1999, Ap	230	10	15	US-10-165-155-1	Sequence 1, Appl
158	15	100.0	10	15	US-10-107-532-2004	Sequence 2004, Ap	231	10	15	US-10-190-162-1	Sequence 1, Appl
159	15	100.0	10	15	US-10-107-532-2025	Sequence 2025, Ap	232	10	15	US-10-224-999A-2225	Sequence 2225, Ap
160	15	100.0	10	15	US-10-107-532-2049	Sequence 2049, Ap	233	10	15	US-10-224-999A-2226	Sequence 2226, Ap
161	15	100.0	10	15	US-10-107-532-2065	Sequence 2065, Ap	234	10	15	US-10-224-999A-2227	Sequence 2227, Ap
	15	100.0	10	15	US-10-107-532-2603	Sequence 2603, Ap		14	14	US-10-224-999A-2228	Sequence 2228, Ap

;; PRIOR FILING DATE: 2001-04-18
;; PRIOR APPLICATION NUMBER: GB 0120084
;; PRIOR FILING DATE: 2001-08-07
;; NUMBER OF SEQ ID NOS: 11
;; SOFTWARE: FASCSQ For Windows Version 4.0
;; SEQ ID NO 11
;; LENGTH: 4
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: 4-mer polypeptide
US-09-998-491-11

Query Match 100.0%; Score 15; DB 10; Length 4;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 2 RER 4

RESULT 4

US-10-357-467-29
; Sequence 29, Application US/10357467
; Publication No. US20030194729A1
; GENERAL INFORMATION:
; APPLICANT: Abogadie, Fe C.
; Cruz, Lourdes J.
; Olivera, Baldomero M.
; Walker, Craig
; Colledge, Clark
; Hilliard, David R.
; Jimenez, Elsie
; TITLE OF INVENTION: Corantokins
; NUMBER OF SEQUENCES: 71
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Rothwell, Figg, Ernst & Manbeck, p.c.
; STREET: 1425 K Street, N.W., Suite 800
; CITY: Washington
; STATE: D.C.
; COUNTRY: USA
; ZIP: 20005
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; FILING DATE: 04-Feb-2003
; APPLICATION NUMBER: US/10/357,467
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 09/142,080
; FILING DATE: 15-MAY-2000
; APPLICATION NUMBER: WO US97/12618
; FILING DATE: 21-JUL-1997
; APPLICATION NUMBER: US 08/684,742
; FILING DATE: 22-JUL-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Ihnen, Jeffrey L.
; REGISTRATION NUMBER: 28,957
; REFERENCE/DOCKET NUMBER: 2314-256.A
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202-783-6040
; TELEFAX: 202-783-6031
; INFORMATION FOR SEQ ID NO: 29:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 4 amino acids
; TYPE: amino acid
; STRANDEDNESS: <Unknown>
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal

;; FEATURE:
;; NAME/KEY: Modified-site
;; LOCATION: 4
;; OTHER INFORMATION: /note= "Xaa is
;; gamma-carboxyglutamic acid"
;; SEQUENCE DESCRIPTION: SEQ ID NO: 29:
US-10-357-467-29

Query Match 100.0%; Score 15; DB 14; Length 4;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 1 RER 3

RESULT 5

US-09-096-749A-59
; Sequence 59, Application US/09096749A
; Patent No. US20020019517A1
; GENERAL INFORMATION:
; APPLICANT: Koieda, Shohei
; TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
; NUMBER OF SEQUENCES: 118
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
; STREET: 121 South Eighth Street, Ste. 1600
; CITY: Minneapolis
; STATE: MN
; COUNTRY: USA
; ZIP: 55402
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 2.0b
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/096,749A
; FILING DATE: June 12, 1998
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Ann S. Viksmans
; REGISTRATION NUMBER: 37,748
; REFERENCE/DOCKET NUMBER: 109.034US1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (612) 373-6900
; TELEFAX: (612) 339-3061
; INFORMATION FOR SEQ ID NO: 59:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
US-09-096-749A-59

Query Match 100.0%; Score 15; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 3 RER 5

RESULT 6

```
US-09-903-412-59
; Sequence 59, Application US/09903412
; Publication No. US20030027319A1
; GENERAL INFORMATION:
; APPLICANT: Koide, Shohel
; TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
; FILE REFERENCE: 109.050051
; CURRENT APPLICATION NUMBER: US/09/903,412
; CURRENT FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: US 60/217,474
; PRIOR FILING DATE: 2000-07-11
; NUMBER OF SEQ ID NOS: 121
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 59
; LENGTH: 5
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: The sequence of the BC loop of clone pLB24.6.
US-09-903-412-59
Query Match 100.0%; Score 15; DB 10; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 7
US-09-998-491-3
; Sequence 3, Application US/09998491
; Publication No. US20030166529A1
; GENERAL INFORMATION:
; APPLICANT: Mileusnic, Radmilla
; APPLICANT: Rose, Stephen Peter Russell
; TITLE OF INVENTION: Polypeptides and their Uses
; FILE REFERENCE: 3578-120
; CURRENT APPLICATION NUMBER: US/09/998,491
; CURRENT FILING DATE: 2001-11-30
; PRIOR APPLICATION NUMBER: GB 0109558.7
; PRIOR FILING DATE: 2001-04-18
; PRIOR APPLICATION NUMBER: GB 0120084
; PRIOR FILING DATE: 2001-08-07
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 3
; LENGTH: 5
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: 5-mer polypeptide
US-09-998-491-3
Query Match 100.0%; Score 15; DB 10; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 1 RER 3

RESULT 8
US-09-998-491-4
; Sequence 4, Application US/09998491
; Publication No. US20030166529A1
; GENERAL INFORMATION:
; APPLICANT: Mileusnic, Radmilla
; APPLICANT: Rose, Stephen Peter Russell
; TITLE OF INVENTION: Polypeptides and their Uses
; FILE REFERENCE: 3578-120
US-09-903-491-4
; CURRENT APPLICATION NUMBER: US/09/998,491
; CURRENT FILING DATE: 2001-11-30
; PRIOR APPLICATION NUMBER: GB 0109558.7
; PRIOR FILING DATE: 2001-04-18
; PRIOR APPLICATION NUMBER: GB 0120084
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4
; LENGTH: 5
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: 5-mer polypeptide
US-09-998-491-4
Query Match 100.0%; Score 15; DB 10; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 9
US-10-174-717A-59
; Sequence 59, Application US/10174717A
; Publication No. US20030108948A1
; APPLICANT: Koide, Shohel
; TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
; NUMBER OF SEQUENCES: 118
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
; STREET: 121 South Eighth Street, St. 1500
; CITY: Minneapolis
; STATE: MN
; COUNTRY: USA
; ZIP: 55402
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: WINDOWS
; SOFTWARE: FastSeq Version 2.0b
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/10/174,717A
; FILING DATE: 18-Jun-2002
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 09/096,749
; FILING DATE: June 12, 1998
; APPLICATION NUMBER: 60/049,410
; FILING DATE: June 12, 1997
; ATTORNEY/AGENT INFORMATION:
; NAME: Ann S. Viksnins
; REGISTRATION NUMBER: 37,748
; REFERENCE/DOCKET NUMBER: 109.034US4
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (612) 373-6900
; TELEFAX: (612) 339-3061
; INFORMATION FOR SEQ ID NO: 59:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHEICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
; SEQUENCE DESCRIPTION: SEQ ID NO: 59:
US-10-174-717A-59
```

Query Match 100.0%; Score 15; DB 14; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 10

US-10-165-155-59
; Sequence 59, Application US/10165155
; Publication No. US20030134386A1
; GENERAL INFORMATION:
; APPLICANT: Koieda, Shohei
; TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
; NUMBER OF SEQUENCES: 118
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
; STREET: 121 South Eighth Street, Ste. 1600
; CITY: Minneapolis
; STATE: MN
; COUNTRY: USA
; ZIP: 55402

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/165,155
FILING DATE: 06-Jun-2002
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/096,749
FILING DATE: June 12, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Ann S. Viksnins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
TELEFAX: (612) 339-3061

INFORMATION FOR SEQ ID NO: 59:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
SEQUENCE DESCRIPTION: SEQ ID NO: 59:
US-10-165-155-59

Query Match 100.0%; Score 15; DB 14; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 11

US-10-190-162-59
; Sequence 59, Application US/10190162
; Publication No. US20030170753A1
; GENERAL INFORMATION:
; APPLICANT: Koieda, Shohei
; TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
; NUMBER OF SEQUENCES: 118

CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, Ste. 1600
CITY: Minneapolis
STATE: MN
COUNTRY: USA
ZIP: 55402

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/190,162
FILING DATE: 03-Jul-2002
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/096,749
FILING DATE: June 12, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Ann S. Viksnins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
TELEFAX: (612) 339-3061

INFORMATION FOR SEQ ID NO: 59:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
SEQUENCE DESCRIPTION: SEQ ID NO: 59:
US-10-190-162-59

Query Match 100.0%; Score 15; DB 14; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 12

US-10-082-747A-60
; Sequence 60, Application US/10082747A
; Publication No. US20030129688A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ballinger, Marcus D.
; APPLICANT: Jones, Jennifer T.
; APPLICANT: Fairbrother, Wayne J.
; APPLICANT: Sliwowski, Mark X.
; APPLICANT: Wells, James A.
; TITLE OF INVENTION: HEREGULIN VARIANTS
; FILE REFERENCE: 402E-476112US
; CURRENT APPLICATION NUMBER: US/10/082,747A
; CURRENT FILING DATE: 2002-09-16
; PRIOR APPLICATION NUMBER: US 09/101,544
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: PCT/US/98/01579
; PRIOR FILING DATE: 1998-02-10
; PRIOR APPLICATION NUMBER: US 08/799,054
; PRIOR FILING DATE: 1997-02-10
; NUMBER OF SEQ ID NOS: 116
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 60
; LENGTH: 6

; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Variant sequence at human heregulin-beta1
; OTHER INFORMATION: residues 183-188
US-10-082-747A-60

Query Match 100.0%; Score 15; DB 14; Length 6;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 2 RER 4

RESULT 13
US-10-315-964A-462
; Sequence 462, Application US/10315964A
; Publication No. US20030148956A1
; GENERAL INFORMATION:
; APPLICANT: The Procter & Gamble Company
; APPLICANT: Isfort, Robert J
; APPLICANT: Mazur, Wieslaw A
; TITLE OF INVENTION: Corticotropin Releasing Factor 2 Receptor Agonists
; FILE REFERENCE: 8847M3
; CURRENT APPLICATION NUMBER: US/10/315,964A
; CURRENT FILING DATE: 2003-04-01
; PRIOR APPLICATION NUMBER: US 60/349,117
; PRIOR FILING DATE: 2002-01-16
; PRIOR APPLICATION NUMBER: US 60/376,337
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: US 60/388,895
; PRIOR FILING DATE: 2002-06-14
; PRIOR APPLICATION NUMBER: US 60/411,988
; PRIOR FILING DATE: 2002-09-19
; NUMBER OF SEQ ID NOS: 530
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 462
; LENGTH: 6
; TYPE: PRT
; ORGANISM: artificial
; FEATURE:
; OTHER INFORMATION: Chemically synthesized artificial peptide
US-10-315-964A-462

Query Match 100.0%; Score 15; DB 14; Length 6;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 14
US-10-315-964A-524
; Sequence 524, Application US/10315964A
; Publication No. US20030148956A1
; GENERAL INFORMATION:
; APPLICANT: The Procter & Gamble Company
; APPLICANT: Isfort, Robert J
; APPLICANT: Mazur, Wieslaw A
; TITLE OF INVENTION: Corticotropin Releasing Factor 2 Receptor Agonists
; FILE REFERENCE: 8847M3
; CURRENT APPLICATION NUMBER: US/10/315,964A
; CURRENT FILING DATE: 2003-04-01
; PRIOR APPLICATION NUMBER: US 60/349,117
; PRIOR FILING DATE: 2002-01-16
; PRIOR APPLICATION NUMBER: US 60/376,337
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: US 60/388,895
; PRIOR FILING DATE: 2002-06-14

; PRIOR APPLICATION NUMBER: US 60/411,988
; PRIOR FILING DATE: 2002-09-19
; NUMBER OF SEQ ID NOS: 530
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 524
; LENGTH: 6
; TYPE: PRT
; ORGANISM: Artificial
; FEATURE:
; OTHER INFORMATION: Chemically synthesized artificial peptide
US-10-315-964A-524

Query Match 100.0%; Score 15; DB 14; Length 6;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 15
US-10-315-964A-525
; Sequence 525, Application US/10315964A
; Publication No. US20030148956A1
; GENERAL INFORMATION:
; APPLICANT: The Procter & Gamble Company
; APPLICANT: Isfort, Robert J
; APPLICANT: Mazur, Wieslaw A
; TITLE OF INVENTION: Corticotropin Releasing Factor 2 Receptor Agonists
; FILE REFERENCE: 8847M3
; CURRENT APPLICATION NUMBER: US/10/315,964A
; CURRENT FILING DATE: 2003-04-01
; PRIOR APPLICATION NUMBER: US 60/349,117
; PRIOR FILING DATE: 2002-01-16
; PRIOR APPLICATION NUMBER: US 60/376,337
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: US 60/388,895
; PRIOR FILING DATE: 2002-06-14
; PRIOR APPLICATION NUMBER: US 60/411,988
; PRIOR FILING DATE: 2002-09-19
; NUMBER OF SEQ ID NOS: 530
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 525
; LENGTH: 6
; TYPE: PRT
; ORGANISM: Artificial
; FEATURE:
; OTHER INFORMATION: Chemically synthesized artificial peptide
US-10-315-964A-525

Query Match 100.0%; Score 15; DB 14; Length 6;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 16
US-10-317-251A-462
; Sequence 462, Application US/10317251A
; Publication No. US20030148957A1
; GENERAL INFORMATION:
; APPLICANT: The Procter & Gamble Company
; APPLICANT: Isfort, Robert J
; APPLICANT: Mazur, Wieslaw A
; TITLE OF INVENTION: Corticotropin Releasing Factor 2 Receptor Agonists
; FILE REFERENCE: 8847M2
; CURRENT APPLICATION NUMBER: US/10/317,251A
; CURRENT FILING DATE: 2002-12-11
; PRIOR APPLICATION NUMBER: US 60/349,117


```
; PRIOR FILING DATE: 2002-01-16
; PRIOR APPLICATION NUMBER: US 60/376,337
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: US 60/388,895
; PRIOR FILING DATE: 2002-06-14
; PRIOR APPLICATION NUMBER: US 60/411,988
; PRIOR FILING DATE: 2002-09-19
; NUMBER OF SEQ ID NOS: 530
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 462
; LENGTH: 6
; TYPE: PRT
; ORGANISM: artificial
; FEATURE:
; OTHER INFORMATION: Chemically synthesized artificial peptide
US-10-317-251A-462

Query Match      100.0%; Score 15; DB 14; Length 6;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 RER 3
Db      3 RER 5

RESULT 17
US-10-317-251A-524
; Sequence 524, Application US/10317251A
; Publication No. US20030148957A1
; GENERAL INFORMATION:
; APPLICANT: The Procter & Gamble Company
; APPLICANT: Isfort, Robert J
; APPLICANT: Mazur, Wieslaw A
; TITLE OF INVENTION: Corticotropin Releasing Factor 2 Receptor Agonists
; FILE REFERENCE: 8847M2
; CURRENT APPLICATION NUMBER: US/10/317,251A
; CURRENT FILING DATE: 2002-12-11
; PRIOR APPLICATION NUMBER: US 60/349,117
; PRIOR FILING DATE: 2002-01-16
; PRIOR APPLICATION NUMBER: US 60/376,337
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: US 60/388,895
; PRIOR FILING DATE: 2002-06-14
; PRIOR APPLICATION NUMBER: US 60/411,988
; NUMBER OF SEQ ID NOS: 530
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 524
; LENGTH: 6
; TYPE: PRT
; ORGANISM: Artificial
; FEATURE:
; OTHER INFORMATION: Chemically synthesized artificial peptide
US-10-317-251A-524

Query Match      100.0%; Score 15; DB 14; Length 6;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 RER 3
Db      3 RER 5

RESULT 18
US-10-317-251A-525
; Sequence 525, Application US/10317251A
; Publication No. US20030148957A1
; GENERAL INFORMATION:
; APPLICANT: The Procter & Gamble Company
; APPLICANT: Isfort, Robert J
; APPLICANT: Mazur, Wieslaw A
```

```
; TITLE OF INVENTION: Corticotropin Releasing Factor 2 Receptor Agonists
; FILE REFERENCE: 8847M2
; CURRENT APPLICATION NUMBER: US/10/317,251A
; CURRENT FILING DATE: 2002-12-11
; PRIOR APPLICATION NUMBER: US 60/349,117
; PRIOR FILING DATE: 2002-01-16
; PRIOR APPLICATION NUMBER: US 60/376,337
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: US 60/388,895
; PRIOR FILING DATE: 2002-06-14
; PRIOR APPLICATION NUMBER: US 60/411,988
; NUMBER OF SEQ ID NOS: 530
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 525
; LENGTH: 6
; TYPE: PRT
; ORGANISM: Artificial
; FEATURE:
; OTHER INFORMATION: Chemically synthesized artificial peptide
US-10-317-251A-525

Query Match      100.0%; Score 15; DB 14; Length 6;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 RER 3
Db      3 RER 5

RESULT 19
US-10-317-252A-462
; Sequence 462, Application US/10317252A
; Publication No. US20030148958A1
; GENERAL INFORMATION:
; APPLICANT: The Procter & Gamble Company
; APPLICANT: Isfort, Robert J
; APPLICANT: Mazur, Wieslaw A
; TITLE OF INVENTION: Corticotropin Releasing Factor 2 Receptor Agonists
; FILE REFERENCE: 8847M
; CURRENT APPLICATION NUMBER: US/10/317,252A
; CURRENT FILING DATE: 2003-03-31
; PRIOR APPLICATION NUMBER: US 60/349,117
; PRIOR FILING DATE: 2002-01-16
; PRIOR APPLICATION NUMBER: US 60/376,337
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: US 60/388,895
; PRIOR FILING DATE: 2002-06-14
; PRIOR APPLICATION NUMBER: US 60/411,988
; NUMBER OF SEQ ID NOS: 530
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 462
; LENGTH: 6
; TYPE: PRT
; ORGANISM: artificial
; FEATURE:
; OTHER INFORMATION: Chemically synthesized artificial peptide
US-10-317-252A-462

Query Match      100.0%; Score 15; DB 14; Length 6;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 RER 3
Db      3 RER 5

RESULT 20
US-10-317-252A-524
; Sequence 524, Application US/10317252A
```


Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 5 RER 7

RESULT 24

US-09-876-904A-495
; Sequence 495, Application US/09876904A
; Publication No. US2003007294A1
; GENERAL INFORMATION:
; APPLICANT: BOULIKAS, TENI
; TITLE OF INVENTION: ENCAPSULATION OF PLASMD DNA (LIPOGENES TM) AND THERAPEUTIC
; TITLE OF INVENTION: AGENTS WITH NUCLEAR LOCALIZATION SIGNAL/FUSOGENIC PEPTIDE
; TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOE COMPLEXES
; FILE REFERENCE: TB-2002.00
; CURRENT APPLICATION NUMBER: US/09/876,904A
; CURRENT FILING DATE: 2001-06-08
; PRIOR FILING DATE: US 60/210,925
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 629
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 495
; LENGTH: 7
; TYPE: PRT
; ORGANISM: Rattus sp.
; FEATURE:
; OTHER INFORMATION: Rat AT-BP1.
US-09-876-904A-495

Query Match 100.0%; Score 15; DB 10; Length 7;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 25

US-09-876-904A-508
; Sequence 508, Application US/09876904A
; Publication No. US2003007294A1
; GENERAL INFORMATION:
; APPLICANT: BOULIKAS, TENI
; TITLE OF INVENTION: ENCAPSULATION OF PLASMD DNA (LIPOGENES TM) AND THERAPEUTIC
; TITLE OF INVENTION: AGENTS WITH NUCLEAR LOCALIZATION SIGNAL/FUSOGENIC PEPTIDE
; TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOE COMPLEXES
; FILE REFERENCE: TB-2002.00
; CURRENT APPLICATION NUMBER: US/09/876,904A
; CURRENT FILING DATE: 2001-06-08
; PRIOR FILING DATE: US 60/210,925
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 629
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 508
; LENGTH: 7
; TYPE: PRT
; ORGANISM: Unknown Organism
; FEATURE:
; OTHER INFORMATION: Description of Unknown Organism: Ig/EBP-1 (immunoglobulin
; OTHER INFORMATION: gene enhancer-binding protein!).
US-09-876-904A-508

Query Match 100.0%; Score 15; DB 10; Length 7;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 5 RER 7

RESULT 26

US-09-972-656-17
; Sequence 17, Application US/09972656
; Publication No. US20030099647A1
; GENERAL INFORMATION:
; APPLICANT: Deshpande, Rajendra
; APPLICANT: Tsai, Mei-Mei
; TITLE OF INVENTION: Fully Human Antibody Fab Fragments with Human Interferon-Gamma
; TITLE OF INVENTION: Neutralizing Activity
; FILE REFERENCE: A-799
; CURRENT APPLICATION NUMBER: US/09/972,656
; CURRENT FILING DATE: 2001-10-05
; NUMBER OF SEQ ID NOS: 135
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 17
; LENGTH: 7
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-972-656-17

Query Match 100.0%; Score 15; DB 10; Length 7;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 27

US-10-232-544-49
; Sequence 49, Application US/10232544
; Publication No. US20030199069A1
; GENERAL INFORMATION:
; APPLICANT: Fuglsang, Claus
; APPLICANT: Okkels, Jens
; APPLICANT: Petersen, Dorte
; APPLICANT: Patkar, Shamkant
; APPLICANT: Thellersen, Marianne
; APPLICANT: Svenden, Allan
; APPLICANT: Borch, Kim
; APPLICANT: Royer, John
; APPLICANT: Kretschmar, Titus
; APPLICANT: Halkier, Torben
; APPLICANT: Vind, Jesper
; APPLICANT: Jorgensen, Steen
; TITLE OF INVENTION: No. US20030199069A1e1 Lipolytic Enzymes
; FILE REFERENCE: 4455.404-US
; CURRENT APPLICATION NUMBER: US/10/232,544
; CURRENT FILING DATE: 2002-08-30
; PRIOR FILING DATE: US/09/007,288
; PRIOR FILING DATE: 2000-01-14
; NUMBER OF SEQ ID NOS: 162
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 49
; LENGTH: 7
; TYPE: PRT
; ORGANISM: Artificial sequence
; FEATURE:
; OTHER INFORMATION: Peptide addition
US-10-232-544-49

Query Match 100.0%; Score 15; DB 14; Length 7;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 4 RER 6

RESULT 28

```

US-09-876-904A-455
; Sequence 455, Application US/09876904A
; Publication No. US20030072794A1
; GENERAL INFORMATION:
; APPLICANT: BOULIKAS, TENI
; TITLE OF INVENTION: ENCAPSULATION OF PLASMID DNA (LIPOGENES TM) AND THERAPEUTIC
; TITLE OF INVENTION: AGENTS WITH NUCLEAR LOCALIZATION SIGNAL/FUSOGENIC PEPTIDE
; TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOME COMPLEXES
; FILE REFERENCE: TB-2002.00
; CURRENT APPLICATION NUMBER: US/09/876,904A
; PRIOR FILING DATE: 2001-06-08
; PRIOR APPLICATION NUMBER: US 60/210,925
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 629
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 455
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Unknown Organism
; FEATURE:
; OTHER INFORMATION: Description of Unknown Organism: C/BBP (CCAAT/enhancer
; OTHER INFORMATION: binding protein).
US-09-876-904A-455

Query Match          100.0%; Score 15; DB 10; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 5 RER 7

RESULT 29
US-09-876-904A-503
; Sequence 503, Application US/09876904A
; Publication No. US20030072794A1
; GENERAL INFORMATION:
; APPLICANT: BOULIKAS, TENI
; TITLE OF INVENTION: ENCAPSULATION OF PLASMID DNA (LIPOGENES TM) AND THERAPEUTIC
; TITLE OF INVENTION: AGENTS WITH NUCLEAR LOCALIZATION SIGNAL/FUSOGENIC PEPTIDE
; TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOME COMPLEXES
; FILE REFERENCE: TB-2002.00
; CURRENT APPLICATION NUMBER: US/09/876,904A
; PRIOR FILING DATE: 2001-06-08
; PRIOR APPLICATION NUMBER: US 60/210,925
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 629
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 503
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Mus sp.
; FEATURE:
; OTHER INFORMATION: Mouse AGP/EBP.
US-09-876-904A-503

Query Match          100.0%; Score 15; DB 10; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 5 RER 7

RESULT 30
US-09-876-904A-506
; Sequence 506, Application US/09876904A
; Publication No. US20030072794A1
; GENERAL INFORMATION:
; APPLICANT: BOULIKAS, TENI
; TITLE OF INVENTION: ENCAPSULATION OF PLASMID DNA (LIPOGENES TM) AND THERAPEUTIC

```

```

; TITLE OF INVENTION: AGENTS WITH NUCLEAR LOCALIZATION SIGNAL/FUSOGENIC PEPTIDE
; TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOME COMPLEXES
; FILE REFERENCE: TB-2002.00
; CURRENT APPLICATION NUMBER: US/09/876,904A
; CURRENT FILING DATE: 2001-06-08
; PRIOR APPLICATION NUMBER: US 60/210,925
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 629
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 506
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Rattus sp.
; FEATURE:
; OTHER INFORMATION: Rat LAP, a 32-kD liver enriched transcriptional
; OTHER INFORMATION: activator, also present in lung, with 71% sequence
; OTHER INFORMATION: similarity to C/BBP.
US-09-876-904A-506

Query Match          100.0%; Score 15; DB 10; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 5 RER 7

RESULT 31
US-09-876-904A-519
; Sequence 519, Application US/09876904A
; Publication No. US20030072794A1
; GENERAL INFORMATION:
; APPLICANT: BOULIKAS, TENI
; TITLE OF INVENTION: ENCAPSULATION OF PLASMID DNA (LIPOGENES TM) AND THERAPEUTIC
; TITLE OF INVENTION: AGENTS WITH NUCLEAR LOCALIZATION SIGNAL/FUSOGENIC PEPTIDE
; TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOME COMPLEXES
; FILE REFERENCE: TB-2002.00
; CURRENT APPLICATION NUMBER: US/09/876,904A
; CURRENT FILING DATE: 2001-06-08
; PRIOR APPLICATION NUMBER: US 60/210,925
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 629
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 519
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Saccharomyces cerevisiae
; FEATURE:
; OTHER INFORMATION: Human NF-IL6 (345 aa).
US-09-876-904A-519

Query Match          100.0%; Score 15; DB 10; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 5 RER 7

RESULT 32
US-09-833-039-39
; Sequence 39, Application US/09833039
; Publication No. US20030175960A1
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfrendschuh, Michael
; TITLE OF INVENTION: Tumor Associated Peptide and Uses Thereof
; FILE REFERENCE: LUD 5622.1
; CURRENT APPLICATION NUMBER: US/09/833,039
; CURRENT FILING DATE: 2001-04-12

```

; PRIOR APPLICATION NUMBER: US 09/409,455
; PRIOR FILING DATE: 1999-09-30
; PRIOR APPLICATION NUMBER: US 09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 129
; SEQ ID NO 39
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-833-039-39

Query Match 100.0%; Score 15; DB 10; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 33
US-09-833-039-64
; Sequence 64, Application US/09833039
; Publication No. US20030175960A1
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfreundschuh, Michael
; TITLE OF INVENTION: Tumor Associated Peptide and Uses Thereof
; FILE REFERENCE: LUD 5622.1
; CURRENT APPLICATION NUMBER: US/09/833,039
; CURRENT FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: US 09/409,455
; PRIOR FILING DATE: 1999-09-30
; PRIOR APPLICATION NUMBER: US 09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 129
; SEQ ID NO 64
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-833-039-64

Query Match 100.0%; Score 15; DB 10; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 34
US-09-833-039-91
; Sequence 91, Application US/09833039
; Publication No. US20030175960A1
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfreundschuh, Michael
; TITLE OF INVENTION: Tumor Associated Peptide and Uses Thereof
; FILE REFERENCE: LUD 5622.1
; CURRENT APPLICATION NUMBER: US/09/833,039
; CURRENT FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: US 09/409,455

; PRIOR FILING DATE: 1999-09-30
; PRIOR APPLICATION NUMBER: US 09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 129
; SEQ ID NO 91
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-833-039-91

Query Match 100.0%; Score 15; DB 10; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 35
US-09-833-039-111
; Sequence 111, Application US/09833039
; Publication No. US20030175960A1
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfreundschuh, Michael
; TITLE OF INVENTION: Tumor Associated Peptide and Uses Thereof
; FILE REFERENCE: LUD 5622.1
; CURRENT APPLICATION NUMBER: US/09/833,039
; CURRENT FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: US 09/409,455
; PRIOR FILING DATE: 1999-09-30
; PRIOR APPLICATION NUMBER: US 09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 129
; SEQ ID NO 111
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-833-039-111

Query Match 100.0%; Score 15; DB 10; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 36
US-10-014-485A-54
; Sequence 54, Application US/10014485A
; Publication No. US20020168684A1
; GENERAL INFORMATION:
; APPLICANT: Cell Signaling Technology, Inc.
; APPLICANT: COMB, Michael J.
; APPLICANT: ZHANG, Hui
; APPLICANT: TAN, Yi
; TITLE OF INVENTION: PRODUCTION OF MOTIF-SPECIFIC AND CONTEXT-INDEPENDENT ANTIBODIES US
; FILE REFERENCE: PEPTIDE LIBRARIES AS ANTIGENS
; FILE REFERENCE: CST-138 CIP2
; CURRENT APPLICATION NUMBER: US/10/014,485A
; CURRENT FILING DATE: 2002-03-18

```
; PRIOR APPLICATION NUMBER: US 09/149,712
; PRIOR FILING DATE: 1998-09-04
; PRIOR APPLICATION NUMBER: US 09/535,364
; PRIOR FILING DATE: 2000-03-24
; NUMBER OF SEQ ID NOS: 145
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 54
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: MOD_RES
; LOCATION: (6)..(6)
; OTHER INFORMATION: PHOSPHORYLATION; threonine at position 6 is phosphorylated
US-10-014-485A-54

Query Match      100.0%; Score 15; DB 13; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 RER 3
Db      1 RER 3

RESULT 37
US-10-174-105A-54
; Sequence 54, Application US/10174105A
; Publication No. US20030068652A1
; GENERAL INFORMATION:
; APPLICANT: Cell Signaling Technology, Inc.
; APPLICANT: ZHANG, Hui
; APPLICANT: COMB, Michael J.
; APPLICANT: TAN, Yi
; TITLE OF INVENTION: POSITIVE IDENTIFICATION OF PHOSPHO-PROTEINS USING MOTIF-SPECIFIC,
; FILE REFERENCE: CST-138 CIP3
; CURRENT APPLICATION NUMBER: US/10/174,105A
; CURRENT FILING DATE: 2002-06-18
; PRIOR APPLICATION NUMBER: US 09/149,712
; PRIOR FILING DATE: 1998-09-04
; PRIOR APPLICATION NUMBER: US 09/535,364
; PRIOR FILING DATE: 2000-03-24
; NUMBER OF SEQ ID NOS: 193
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 54
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic Peptide
; NAME/KEY: MOD_RES
; LOCATION: (6)..(6)
; OTHER INFORMATION: PHOSPHORYLATION; threonine at position 6 is phosphorylated
US-10-174-105A-54

Query Match      100.0%; Score 15; DB 14; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 RER 3
Db      1 RER 3

RESULT 38
US-10-224-999A-2180
; Sequence 2180, Application US/10224999A
; Publication No. US20030171318A1
; GENERAL INFORMATION:
; APPLICANT: Myriad Genetics, Inc.
; APPLICANT: Morham, Scott
```

```
; APPLICANT: Zavitz, Kenton
; APPLICANT: Hobden, Adrian
; TITLE OF INVENTION: Composition and Method for Treating Viral Infection
; FILE REFERENCE: 5004.01
; CURRENT APPLICATION NUMBER: US/10/224,999A
; CURRENT FILING DATE: 2003-03-03
; PRIOR APPLICATION NUMBER: US 60/313,695
; PRIOR FILING DATE: 2001-08-20
; NUMBER OF SEQ ID NOS: 3484
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 2180
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Rubella virus
US-10-224-999A-2180

Query Match      100.0%; Score 15; DB 14; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 RER 3
Db      2 RER 4

RESULT 39
US-10-224-999A-2181
; Sequence 2181, Application US/10224999A
; Publication No. US20030171318A1
; GENERAL INFORMATION:
; APPLICANT: Myriad Genetics, Inc.
; APPLICANT: Morham, Scott
; APPLICANT: Zavitz, Kenton
; APPLICANT: Hobden, Adrian
; TITLE OF INVENTION: Composition and Method for Treating Viral Infection
; FILE REFERENCE: 5004.01
; CURRENT APPLICATION NUMBER: US/10/224,999A
; CURRENT FILING DATE: 2003-03-03
; PRIOR APPLICATION NUMBER: US 60/313,695
; PRIOR FILING DATE: 2001-08-20
; NUMBER OF SEQ ID NOS: 3484
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 2181
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Rubella virus
US-10-224-999A-2181

Query Match      100.0%; Score 15; DB 14; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 RER 3
Db      1 RER 3

RESULT 40
US-10-177-277-39
; Sequence 39, Application US/10177277
; Publication No. US2003018544A1
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfreundschuh, Michael
; APPLICANT: Ramensee, Hans Georg
; APPLICANT: Stevanovic, Stefan
; TITLE OF INVENTION: Method for Determining Presence of Cancer In a Sample By Determini
; TITLE OF INVENTION: Expression of an SSX Gene, Peptides Derived From Said SSX Gene ar
; TITLE OF INVENTION: Gene, and Uses Thereof
; FILE REFERENCE: LUD 5556.1
; CURRENT APPLICATION NUMBER: US/10/177,277
; CURRENT FILING DATE: 2002-06-21
```

; FILE REFERENCE: LUD 5556.1
; CURRENT APPLICATION NUMBER: US/09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 132
; SEQ ID NO 39
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-177-277-39

Query Match 100.0%; Score 15; DB 14; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 41

US-10-177-277-64
; Sequence 64, Application US/10177277
; Publication No. US20030185844A1

; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfreundschuh, Michael
; APPLICANT: Ramensee, Hans Georg
; APPLICANT: Stevanovic, Stefan
; TITLE OF INVENTION: Method for Determining Presence of Cancer In a Sample By Determining Presence of an SSX Gene, Peptides Derived From Said SSX Gene and Uses Thereof
; FILE REFERENCE: LUD 5556.1
; CURRENT APPLICATION NUMBER: US/10/177,277
; PRIOR FILING DATE: 2002-06-21
; PRIOR APPLICATION NUMBER: US/09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 132
; SEQ ID NO 64
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-177-277-64

Query Match 100.0%; Score 15; DB 14; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 42

US-10-177-277-91
; Sequence 91, Application US/10177277
; Publication No. US20030185844A1

; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfreundschuh, Michael
; APPLICANT: Ramensee, Hans Georg
; APPLICANT: Stevanovic, Stefan
; TITLE OF INVENTION: Method for Determining Presence of Cancer In a Sample By Determining Presence of an SSX Gene, Peptides Derived From Said SSX Gene and Uses Thereof
; FILE REFERENCE: LUD 5556.1
; CURRENT APPLICATION NUMBER: US/10/177,277
; PRIOR FILING DATE: 2002-06-21
; PRIOR APPLICATION NUMBER: US/09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 132
; SEQ ID NO 91
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-177-277-91

Query Match 100.0%; Score 15; DB 14; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

; FILE REFERENCE: LUD 5556.1
; CURRENT APPLICATION NUMBER: US/10/177,277
; PRIOR FILING DATE: 2002-06-21
; PRIOR APPLICATION NUMBER: US/09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 132
; SEQ ID NO 91
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-177-277-91

Query Match 100.0%; Score 15; DB 14; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 43

US-10-177-277-111
; Sequence 111, Application US/10177277
; Publication No. US20030185844A1

; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfreundschuh, Michael
; APPLICANT: Ramensee, Hans Georg
; APPLICANT: Stevanovic, Stefan
; TITLE OF INVENTION: Method for Determining Presence of Cancer In a Sample By Determining Presence of an SSX Gene, Peptides Derived From Said SSX Gene and Uses Thereof
; FILE REFERENCE: LUD 5556.1
; CURRENT APPLICATION NUMBER: US/10/177,277
; PRIOR FILING DATE: 2002-06-21
; PRIOR APPLICATION NUMBER: US/09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 132
; SEQ ID NO 111
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-177-277-111

Query Match 100.0%; Score 15; DB 14; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 44

US-10-376-121A-79
; Sequence 79, Application US/10376121A
; Publication No. US20030216544A1

; GENERAL INFORMATION:
; APPLICANT: Harley, John
; TITLE OF INVENTION: METHODS AND REAGENTS FOR DIAGNOSIS OF AUTOANTIBODIES
; NUMBER OF SEQUENCES: 218
; CORRESPONDENCE ADDRESS:
US-10-376-121A-79

Query Match 100.0%; Score 15; DB 14; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

ADDRESSEE: Patrea L. Pabst
STREET: Suite 2000, 1201 West Peachtree Street, N.E.
CITY: Atlanta
STATE: GA
COUNTRY: USA
ZIP: 30309-3400
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/376,121A
FILING DATE: 27-Mar-2003
CLASSIFICATION: <Unknown>
PRIORITY APPLICATION DATA:
APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: OMRF114CIP(2)DIV(2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404)-817-8473
TELEFAX: (404)-817-8588
INFORMATION FOR SEQ ID NO: 79:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
FEATURE:
NAME/KEY: Binding-site
LOCATION: 1..8
SEQUENCE DESCRIPTION: SEQ ID NO: 79:
US-10-376-121A-79
Query Match 100.0%; Score 15; DB 15; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
DB 5 RER 7
RESULT 45
US-10-376-121A-80
; Sequence 80, Application US/10376121A
; Publication No. US200302165441
; GENERAL INFORMATION:
; APPLICANT: Harley, John
; TITLE OF INVENTION: METHODS AND REAGENTS FOR DIAGNOSIS OF
; AUTOANTIBODIES
; NUMBER OF SEQUENCES: 218
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Patrea L. Pabst
; STREET: Suite 2000, 1201 West Peachtree Street, N.E.
; CITY: Atlanta
; STATE: GA
; COUNTRY: USA
; ZIP: 30309-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/376,121A
FILING DATE: 27-Mar-2003
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/867,819
FILING DATE: April 13, 1992
APPLICATION NUMBER: 07/648,205
FILING DATE: January 31, 1991
APPLICATION NUMBER: 07/472,947
FILING DATE: January 31, 1990
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: OMRF114CIP(2)DIV(2)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404)-817-8473
TELEFAX: (404)-817-8588
INFORMATION FOR SEQ ID NO: 80:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
FEATURE:
NAME/KEY: Binding-site
LOCATION: 1..8
SEQUENCE DESCRIPTION: SEQ ID NO: 80:
US-10-376-121A-80
Query Match 100.0%; Score 15; DB 15; Length 8;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
DB 2 RER 4
RESULT 46
US-09-809-638-516
; Sequence 516, Application US/09809638
; Publication No. US20030059895A1
; GENERAL INFORMATION:
; APPLICANT: Mary Faris
; APPLICANT: Pia M. Challita-Bid
; APPLICANT: Steve Chappell Mitchell
; APPLICANT: Daniel E.H. Afar
; APPLICANT: Arthur B. Raitano
; APPLICANT: Aya Jakobovits
; TITLE OF INVENTION: 125P5C8: A TISSUE SPECIFIC PROTEIN
; FILE REFERENCE: 129.35US01
; CURRENT APPLICATION NUMBER: US/09/809,638
; CURRENT FILING DATE: 2001-03-14
; NUMBER OF SEQ ID NOS: 746
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 516
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-809-638-516
Query Match 100.0%; Score 15; DB 10; Length 9;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
DB 3 RER 5
RESULT 47

US-09-809-638-621
; Sequence 621, Application US/09809638
; Publication No. US20030059895A1
; GENERAL INFORMATION:

; APPLICANT: Mary Faris
; APPLICANT: Pia M. Challita-Eid
; APPLICANT: Steve Chappell Mitchell
; APPLICANT: Daniel E.H. Afar
; APPLICANT: Arthur B. Raitano
; APPLICANT: Aya Jakobovits
; TITLE OF INVENTION: 125P5C8: A TISSUE SPECIFIC PROTEIN
; FILE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCERS
; FILE REFERENCE: 129.35US01
; CURRENT APPLICATION NUMBER: US/09/809,638
; CURRENT FILING DATE: 2001-03-14
; NUMBER OF SEQ ID NOS: 746
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 621
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-809-638-621

Query Match 100.0%; Score 15; DB 10; Length 9;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 48

US-09-876-904A-379
; Sequence 379, Application US/09876904A
; Publication No. US20030072794A1
; GENERAL INFORMATION:

; APPLICANT: BOULIKAS, TENI
; TITLE OF INVENTION: ENCAPSULATION OF PLASMID DNA (LIPOGENES TM) AND THERAPEUTIC
; TITLE OF INVENTION: AGENTS WITH NUCLEAR LOCALIZATION SIGNAL/FUSOGENIC PEPTIDE
; TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOOME COMPLEXES
; FILE REFERENCE: TB-2002.00
; CURRENT APPLICATION NUMBER: US/09/876,904A
; CURRENT FILING DATE: 2001-06-08
; PRIOR APPLICATION NUMBER: US 60/210,925
; PRIOR FILING DATE: 2000-06-09
; NUMBER OF SEQ ID NOS: 629
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 379
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: Human thyroid hormone receptor alpha (c-erbA-1
US-09-876-904A-379

Query Match 100.0%; Score 15; DB 10; Length 9;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 2 RER 4

RESULT 49

US-09-932-165-1036
; Sequence 1036, Application US/09932165
; Publication No. US20030134784A1
; GENERAL INFORMATION:
; APPLICANT: RAITANG, ARTHUR
; APPLICANT: CHALLITA-EID, PIA M.

; APPLICANT: FARIS, MARY
; APPLICANT: SAFFRAN, DOUGLAS
; APPLICANT: AFAR, DANIEL
; APPLICANT: LEVIN, ELANA
; APPLICANT: HUBERT, RENE
; APPLICANT: GE, WANGMAO
; APPLICANT: JAKOBOVITS, AYA
; TITLE OF INVENTION: NUCLEIC ACIDS AND CORRESPONDING PROTEINS ENTITLED
; TITLE OF INVENTION: 83P2H3 AND Cat-F2E11 USEFUL IN TREATMENT AND
; TITLE OF INVENTION: DETECTION OF CANCER
; FILE REFERENCE: 51158-20014.00
; CURRENT APPLICATION NUMBER: US/09/932,165
; CURRENT FILING DATE: 2001-08-17
; PRIOR APPLICATION NUMBER: 60/226,329
; PRIOR FILING DATE: 2000-08-17
; NUMBER OF SEQ ID NOS: 1508
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1036
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Peptide motif
US-09-932-165-1036

Query Match 100.0%; Score 15; DB 10; Length 9;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 50

US-09-833-039-41
; Sequence 41, Application US/09833039
; Publication No. US20030175960A1
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfrendtschuh, Michael
; TITLE OF INVENTION: Tumor Associated Peptide and Uses Thereof
; FILE REFERENCE: LUD 5622.1
; CURRENT APPLICATION NUMBER: US/09/833,039
; CURRENT FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: US 09/409,455
; PRIOR FILING DATE: 1999-09-30
; PRIOR APPLICATION NUMBER: US 09/344,040
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 129
; SEQ ID NO 41
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-833-039-41

Query Match 100.0%; Score 15; DB 10; Length 9;
Best Local Similarity 100.0%; Pred. No. 7.1e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 1 RER 3

Search completed: March 5, 2004, 16:12:26
Job time : 33 secs

GenCore version 5.1.6
Copyright (c) 1993 - 2004 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: March 5, 2004, 16:06:22 ; Search time 21 Seconds
(without alignments)
13.742 Million cell updates/sec

Title: US-09-998-491-9

Perfect score: 15

Sequence: 1 RER 3

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 28356 seqs, 96191526 residues

Total number of hits satisfying chosen parameters: 5

Minimum DB seq length: 0

Maximum DB seq length: 20

Post-processing: Minimum Match 100%

Maximum Match 100%

Listing first 1000 summaries

Database : PIR-78:*

1: PIR1.*

2: PIR2.*

3: PIR3.*

4: PIR4.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	15	100.0	12	T44420	hypothetical prote
2	15	100.0	13	PH1595	Ig H chain V-D-J r
3	15	100.0	15	S57584	T cell receptor V-
4	15	100.0	16	PH1475	T-cell receptor be
5	15	100.0	18	S54270	GATA-2 protein - A

ALIGNMENTS

RESULT 1

T44420

hypothetical protein [imported] - Bacillus stearothermophilus (fragment)

C:Species: Bacillus stearothermophilus

C>Date: 21-Jan-2000 #sequence_revision 21-Jan-2000 #text_change 21-Jan-2000

C:Accession: T44420

R:Vlaskova, H.; Krasny, L.; Fucik, V.; Jonak, J.

submitted to the EMBL Data Library, September 1997

A:Description: The pyrAB gene coding for the large subunit of carbamoylphosphate synthet

A:Reference number: Z22760

A:Accession: T44420

A>Status: preliminary; translated from GB/EMBL/DBJ

A:Molecule type: DNA

A:Residues: 1-12 <VLA>

A:Cross-references: EMBL:AJ001805; PIDN:CAA05021.1

A:Experimental source: strain CCM 2184

C:Genetics:

A>Note: ORF2

Query Match 100.0%; Score 15; DB 2; Length 12;
Best Local Similarity 100.0%; Pred. No. 3e+02;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3

Db 4 RER 6

RESULT 2

PH1595

Ig H chain V-D-J region (wild-type clone 150) - mouse (fragment)

C:Species: Mus musculus (house mouse)

C>Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999

C:Accession: PH1595

R:Levinson, D.A.; Campos-Torres, J.; Leder, P.

J. Exp. Med. 178, 317-329, 1993

A>Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice

A:Reference number: PH1580; MUID:93301609; PMID:8315387

A:Accession: PH1595

A:Molecule type: DNA

A:Residues: 1-13 <LEV>

A:Experimental source: bone marrow pre-B lymphocyte

C:Keywords: immunoglobulin

Query Match 100.0%; Score 15; DB 2; Length 13;

Best Local Similarity 100.0%; Pred. No. 3.2e+02;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3

Db 3 RER 5

RESULT 3

S57584

T cell receptor V-D-J junctional alpha chain region - human (fragment)

C:Species: Homo sapiens (man)

C>Date: 19-Oct-1995 #sequence_revision 17-Nov-1995 #text_change 05-Nov-1999

C:Accession: S57584

R:Burrows, S.R.; Silins, S.L.; Moss, D.J.; Khanna, R.; Misko, I.S.; Arguet, V.P.

submitted to the EMBL Data Library, June 1995

A:Description: T cell receptor repertoire for a viral epitope in humans is diversified by

A:Reference number: S57494

A:Accession: S57584

A>Status: preliminary

A:Molecule type: mRNA

A:Residues: 1-15 <BUR>

A:Cross-references: EMBL:Z49556; NID:g887466; PIDN:CAA90227.1; PMID:9887467

C:Keywords: T-cell receptor

Query Match 100.0%; Score 15; DB 2; Length 15;

Best Local Similarity 100.0%; Pred. No. 3.7e+02;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3

Db 6 RER 8

RESULT 4

PH1475

T-cell receptor beta chain (clone 223/5) - mouse (fragment)

C:Species: Mus musculus (house mouse)

C>Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 11-Apr-1995

C:Accession: PH1475

R:Casanova, J.L.; Martinon, F.; Gournier, H.; Barra, C.; Pannetier, C.; Regnault, A.; Ko

J. Exp. Med. 177, 811-820, 1993

A>Title: T cell receptor selection by and recognition of two class I major histocompatibi

A:Reference number: PH1430; MUID:93171821; PMID:8436911

A:Accession: PH1475

A:Molecule type: mRNA

A:Residues: 1-16 <CAS>

A:Experimental source: cytolytic T-lymphocyte
C:Superfamily: immunoglobulin homology
C:Keywords: receptor; T-cell

Query Match 100.0%; Score 15; DB 2; Length 16;
Best Local Similarity 100.0%; Pred. No. 3.9e+02;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 10 RER 12

RESULT 5

S54270
GATA-2 protein - African clawed frog
C:Species: Xenopus laevis (African Clawed frog)
C>Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 03-Nov-1995
C:Accession: S54270
R:Brewer, A.C.; Guille, M.J.; Fear, D.J.; Partington, G.A.; Patient, R.K.
EMBO J. 14, 757-766, 1995
A:Title: Nuclear translocation of a maternal CCAAT factor at the start of gastrulation
A:Reference number: S54270; MUID:95188880; PMID:7862979
A:Accession: S54270
A>Status: preliminary
A:Molecule type: DNA
A:Residues: 1-18 <BRE>

Query Match 100.0%; Score 15; DB 2; Length 18;
Best Local Similarity 100.0%; Pred. No. 4.4e+02;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 9 RER 11

Search completed: March 5, 2004, 16:11:05
Job time : 22 secs

GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.

OM protein - protein search, using sw model

Run on: March 5, 2004, 16:04:07 ; Search time 11 Seconds
(without alignments)
14.201 Million cell updates/sec

Title: US-09-998-491-9
Perfect score: 15
Sequence: 1 RER 3

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 141681 seqs, 52070155 residues

Total number of hits satisfying chosen parameters: 0

Minimum DB seq length: 0
Maximum DB seq length: 20

Post-processing: Minimum Match 100%
Maximum Match 100%
Listing first 1000 summaries

Database : SwissProt_42:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Score	Match	Length	DB ID	Description
-----	-----	-----	-----	-----	-----

No matches found

Search completed: March 5, 2004, 16:09:40
Job time : 11 secs

GenCore version 5.1.6
Copyright (c) 1993 - 2004 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: March 5, 2004, 16:05:07 ; Search time 38 Seconds
(without alignments)
24.909 Million cell updates/sec

Title: US-09-998-491-9
Perfect score: 15
Sequence: 1 RER 3

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1017041 seqs, 315518202 residues

Total number of hits satisfying chosen parameters: 9

Minimum DB seq length: 0
Maximum DB seq length: 20

Post-processing: Minimum Match 100%
Maximum Match 100%
Listing first 1000 summaries

Database :

SPTREMBL 25: *
1: sp archaea: *
2: sp bacteria: *
3: sp fungi: *
4: sp human: *
5: sp invertebrate: *
6: sp mammal: *
7: sp mhc: *
8: sp organelle: *
9: sp phage: *
10: sp plant: *
11: sp rodent: *
12: sp virus: *
13: sp vertebrate: *
14: sp unclassified: *
15: sp virus: *
16: sp bacterioph: *
17: sp archaeop: *

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	15	100.0	11	Q9S8X4	Q9S8X4 glycine max
2	15	100.0	12	O50303	O50303 bacillus st
3	15	100.0	15	O35411	O35411 mus musculus
4	15	100.0	16	Q9UD21	Q9UD21 homo sapien
5	15	100.0	16	Q9T2R8	Q9T2R8 solanum tub
6	15	100.0	17	Q9CEX8	Q9CEX8 human immun
7	15	100.0	17	Q9CEX9	Q9CEX9 human immun
8	15	100.0	18	Q9GNH1	Q9GNH1 macaca mula
9	15	100.0	20	Q8NEF1	Q8NEF1 homo sapien

ALIGNMENTS

RESULT 1
Q9S8X4

```
ID Q9S8X4 PRELIMINARY; PRT; 11 AA.
AC Q9S8X4;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT 01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
DE Vegetative storage protein 94 peptide 3, VSP94=LIPXYGENASE
DE (fragment).
OS Glycine max (Soybean).
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
OC Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
OC eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae; Glycine.
OX NCBI_TaxID=3847;
RN [1]
RP SEQUENCE.
RX MEDLINE=92361246; PubMed=1822994;
RA Tranbarger T.J., Franceschi V.R., Hildebrand D.F., Grimes H.D.;
RT "The soybean 94-kilodalton vegetative storage protein is a
RT lipoygenase that is localized in paraveinal mesophyll cell
RT vacuoles."
RL Plant Cell 3:973-987(1991).
FT NON_TER 1 1
FT NON_TER 11 11
SQ SEQUENCE 11 AA; 1366 MW; 9B337C3CDD9CB1A CRC64;

Query Match 100.0%; Score 15; DB 10; Length 11;
Best Local Similarity 100.0%; Pred. No. 1.5e+03;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 8 RER 10

RESULT 2
O50303 PRELIMINARY; PRT; 12 AA.
AC O50303;
DT 01-JUN-1998 (TrEMBLrel. 06, Created)
DT 01-JUN-1998 (TrEMBLrel. 06, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein (Fragment).
OS Bacillus stearothermophilus.
OC Bacteria; Firmicutes; Bacillales; Bacillaceae; Geobacillus.
OX NCBI_TaxID=1422;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=COM 2184;
RX MEDLINE=20194845; PubMed=10732707;
RA Vlasikova H., Krasny L., Fucik V., Jonak J.;
RT "The pyrab Gene Coding for the Large Subunit of Carbamoylphosphate
RT Synthetase from Bacillus stearothermophilus: Molecular cloning and
RT Functional Characterization."
RL Folia Biol. (Praha) 44:163-172(1998).
DR EMBL; AJ001805; CAA05021.1; -.
DR PIR; T4420; T4420.
KW Hypothetical protein.
FT NON_TER 12 12
SQ SEQUENCE 12 AA; 1379 MW; 70087CB0E8A6840B CRC64;

Query Match 100.0%; Score 15; DB 2; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.6e+03;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 4 RER 6

RESULT 3
O35411 PRELIMINARY; PRT; 15 AA.
ID O35411
AC O35411;
DT 01-JAN-1998 (TrEMBLrel. 05, Created)
```

```
DT 01-JAN-1998 (TrEMBLrel. 05, last sequence update)
DT 01-DEC-2001 (TrEMBLrel. 19, last annotation update)
DE Beta III spectrin (Fragment).
GN SPNB3.
OS Mus musculus (Mouse).
OC Eukaryota; Eutheria; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
CX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=C57BL/6J; TISSUE=Embryo;
RX MEDLINE=99045634; PubMed=9826670;
RA Stankewich M.C., Tse W.T., Peters L.L., Ch'ng Y., John K.M.,
RA Stabach P.R., Devarajan P., Morrow J.S., Lux S.E.;
RT "A widely expressed betaIII spectrin associated with Golgi and
RT cytoplasmic vesicles.";
RL Proc. Natl. Acad. Sci. U.S.A. 95:14158-14163 (1998).
DR EMBL; AF026489; AAC79505.1; -.
DR MGD; MGI:1313261; Spnb3.
FT NON_TER 1
SQ SEQUENCE 15 AA; 2029 MW; CAF6B165F69F1AA8 CRC64;

Query Match 100.0%; Score 15; DB 11; Length 15;
Best Local Similarity 100.0%; Pred. No. 2e+03;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 2 RER 4

RESULT 4
Q9UD21
ID Q9UD21 PRELIMINARY; PRT; 16 AA.
AC Q9UD21;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, last sequence update)
DT 01-JUN-2000 (TrEMBLrel. 14, last annotation update)
DE Cyclin E-L (Fragment).
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.
CX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC MEDLINE=95257942; PubMed=7739542;
RA Ohtsubo M., Theodoras A.M., Schumacher J., Roberts J.M., Pagano M.;
RA "Human cyclin E, a nuclear protein essential for the G1-to-S phase
RT transition.";
RL Mol. Cell. Biol. 15:2612-2624 (1995).
DR EMBL; AF04368.1; -.
FT NON_TER 1
SQ SEQUENCE 16 AA; 2089 MW; 777EFC69CA45E29C CRC64;

Query Match 100.0%; Score 15; DB 4; Length 16;
Best Local Similarity 100.0%; Pred. No. 2.1e+03;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 5
Q9T2R8
ID Q9T2R8 PRELIMINARY; PRT; 16 AA.
AC Q9T2R8;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, last annotation update)
DE Cytochrome-c reductase 53 kDa subunit (EC 1.10.2.2) (Fragment).
OS Solanum tuberosum (Potato).
OC Mitochondrion.
OC Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
OC Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
```

```
OC lamids; Solanales; Solanaceae; Solanum.
OX NCBI_TaxID=4113;
RN [1]
RP SEQUENCE.
RX MEDLINE=94198758; PubMed=7764624;
RA Braun H.P., Kruft V., Schmitz U.K.;
RA Planta 193:199-106 (1994).
RL GO; GO:0008121; F:ubiquinol-cytochrome-c reductase activity; IEA.
DR EMBL; AF04367.1; -.
FT NON_TER 1
SQ SEQUENCE 16 AA; 2116 MW; 91C55A205F04C82 CRC64;

Query Match 100.0%; Score 15; DB 8; Length 16;
Best Local Similarity 100.0%; Pred. No. 2.1e+03;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 9 RER 11

RESULT 6
Q9QEX8
ID Q9QEX8 PRELIMINARY; PRT; 17 AA.
AC Q9QEX8;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, last sequence update)
DT 01-OCT-2002 (TrEMBLrel. 22, last annotation update)
DE Nef protein (Fragment).
GN NEF.
OS Human immunodeficiency virus 1.
OC Viruses; Retroviridae; Retroviridae; Lentivirus.
CX NCBI_TaxID=11676;
RN [1]
RP SEQUENCE FROM N.A.
RC MEDLINE=21103026; PubMed=11170057;
RA Lin H.J., Sivak E.B., Lauder I.J., Hollinger F.B.;
RA "Long-term culture of human immunodeficiency virus type 1 resulting in
RT loss of glycosylation sites.";
RL J. Med. Virol. 63:197-202 (2001).
DR EMBL; AF178662; AAF04368.1; -.
FT NON_TER 1
FT NON_TER 17
SQ SEQUENCE 17 AA; 2032 MW; 919FC3C6F3515653 CRC64;

Query Match 100.0%; Score 15; DB 15; Length 17;
Best Local Similarity 100.0%; Pred. No. 2.2e+03;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 14 RER 16

RESULT 7
Q9QEX9
ID Q9QEX9 PRELIMINARY; PRT; 17 AA.
AC Q9QEX9;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, last sequence update)
DT 01-OCT-2002 (TrEMBLrel. 22, last annotation update)
DE Nef protein (Fragment).
GN NEF.
OS Human immunodeficiency virus 1.
OC Viruses; Retroviridae; Retroviridae; Lentivirus.
CX NCBI_TaxID=11676;
RN [1]
RP SEQUENCE FROM N.A.
RC MEDLINE=21103026; PubMed=11170057;
RA Lin H.J., Sivak E.B., Lauder I.J., Hollinger F.B.;
RA "Long-term culture of human immunodeficiency virus type 1 resulting in
RT loss of glycosylation sites.";
RL J. Med. Virol. 63:197-202 (2001).
DR EMBL; AF178663; AAF04367.1; -.
FT NON_TER 1
FT NON_TER 1
```

```

FT  NON TER      17
SQ  SEQUENCE 17 AA; 1960 MW; 9315C3C6F3515653 CRC64;

Query Match      100.0%; Score 15; DB 15; Length 17;
Best Local Similarity 100.0%; Pred. No. 2.2e+03;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY  1 RER 3
Db  14 RER 16

RESULT 8
Q8GNEI
ID  Q8GNEI      PRELIMINARY; PRT; 18 AA.
AC  Q8GNEI;
DT  01-MAR-2001 (TREMBLrel. 16, Created)
DT  01-MAR-2001 (TREMBLrel. 16, Last sequence update)
DT  01-MAR-2001 (TREMBLrel. 16, Last annotation update)
DE  Matrix Gla protein (Fragment).
GN  MGP.
OS  Macaca mulatta (Rhesus macaque).
OC  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC  Mammalia; Eutheria; Primates; Catarrhini; Cercopithecoidea;
OC  Cercopithecinae; Macaca.
OX  NCBI_TaxID=9544;
RN  [1]
RP  SEQUENCE FROM N.A.
RC  TISSUE=Carotid artery;
RA  Wu K.-J., Yee A., Zhu N.L., Gordon E.M., Hall F.L.;
RT  "Characterization of differential gene expression in monkey arterial
RT  neointima following balloon catheter injury.";
RL  Submitted (JUN-1999) to the EMBL/GenBank/DBJ databases.
DR  EMBL; AF162477; AAF98709.1; -.
FT  NON TER      1
SQ  SEQUENCE 18 AA; 2255 MW; FB4F252C395E5DB1 CRC64;

Query Match      100.0%; Score 15; DB 6; Length 18;
Best Local Similarity 100.0%; Pred. No. 2.3e+03;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY  1 RER 3
Db  12 RER 14

RESULT 9
Q8NEEI
ID  Q8NEEI      PRELIMINARY; PRT; 20 AA.
AC  Q8NEEI;
DT  01-OCT-2002 (TREMBLrel. 22, Created)
DT  01-OCT-2002 (TREMBLrel. 22, Last sequence update)
DT  01-OCT-2002 (TREMBLrel. 22, Last annotation update)
DE  Hypothetical protein (Fragment).
OS  Homo sapiens (Human).
OC  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC  Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX  NCBI_TaxID=9606;
RN  [1]
RP  SEQUENCE FROM N.A.
RC  TISSUE=Testis;
RA  Strausberg R.;
RL  Submitted (JUN-2002) to the EMBL/GenBank/DBJ databases.
DR  EMBL; BC031872; AAH31872.1; -.
KW  Hypothetical protein.
FT  NON TER      1
SQ  SEQUENCE 20 AA; 2218 MW; 8C8A0AD4BF387987 CRC64;

Query Match      100.0%; Score 15; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 2.6e+03;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY  1 RER 3

```

Db 1 RER 3

Search completed: March 5, 2004, 16:10:32
Job time : 39 secs

99	15	100.0	9	2	RAY20470	Aay20470	Human mic	172	15	100.0	9	6	ABU77246	Novel pro
100	15	100.0	9	2	AAW1588	Aaw1588	Melanoma	173	15	100.0	9	6	ABJ38075	Human cye
101	15	100.0	9	2	RAY01011	Aay01011	Bacterial	174	15	100.0	9	6	ABJ20567	162P1E6 c
102	15	100.0	9	3	RAY76726	Aay76726	SSX-2 H1A	175	15	100.0	9	6	ABJ20602	162P1E6 c
103	15	100.0	9	3	RAY76745	Aay76745	SSX-1 H1A	176	15	100.0	9	6	ABJ21965	162P1E6 c
104	15	100.0	9	3	RAY76735	Aay76735	SSX-2 H1A	177	15	100.0	9	6	ABJ21994	162P1E6 c
105	15	100.0	9	3	RAY78491	Aay78491	SSX-5 der	178	15	100.0	9	6	ABJ24087	162P1E6 c
106	15	100.0	9	3	RAY79727	Aay79727	SSX-2 der	179	15	100.0	9	6	ABJ24088	162P1E6 c
107	15	100.0	9	3	RAY79701	Aay79701	SSX-4 der	180	15	100.0	9	6	ABJ20598	162P1E6 c
108	15	100.0	9	3	RAY79708	Aay79708	SSX-2 der	181	15	100.0	9	6	ABJ21980	162P1E6 c
109	15	100.0	9	3	RAY79745	Aay79745	SSX-1 der	182	15	100.0	9	6	ABJ22659	162P1E6 c
110	15	100.0	9	4	RAM99326	Aam99326	Vaccine r	183	15	100.0	9	6	ABJ24748	162P1E6 c
111	15	100.0	9	4	RAM98902	Aam98902	Vaccine r	184	15	100.0	9	6	ABJ24743	162P1E6 c
112	15	100.0	9	4	ABP15116	Abp15116	HIV A03 s	185	15	100.0	9	6	ABJ22691	162P1E6 c
113	15	100.0	9	4	ABP20861	Abp20861	HIV A03 m	186	15	100.0	9	6	ABJ23377	162P1E6 c
114	15	100.0	9	4	ABP22956	Abp22956	HIV A11 m	187	15	100.0	9	6	ABJ21271	162P1E6 c
115	15	100.0	9	4	ABP16400	Abp16400	HIV A24 s	188	15	100.0	9	6	ABJ21985	162P1E6 c
116	15	100.0	9	4	ABP22268	Abp22268	HIV A03 m	189	15	100.0	9	6	ABJ23386	162P1E6 c
117	15	100.0	9	4	ABP14244	Abp14244	HIV A02 s	190	15	100.0	9	6	ABJ20591	162P1E6 c
118	15	100.0	9	4	ABP24326	Abp24326	HIV A24 m	191	15	100.0	9	6	ABJ21265	162P1E6 c
119	15	100.0	9	4	ABP22259	Abp22259	HIV A03 m	192	15	100.0	9	6	ABJ23369	162P1E6 c
120	15	100.0	9	4	ABP24024	Abp24024	HIV A11 m	193	15	100.0	9	6	ABJ21286	162P1E6 c
121	15	100.0	9	4	AAAM49825	Aam49825	Human D40	194	15	100.0	9	6	ABJ24785	162P1E6 c
122	15	100.0	9	5	ABP74615	Abp74615	Transcrip	195	15	100.0	9	6	ABJ24754	162P1E6 c
123	15	100.0	9	5	AAE18744	Aae18744	Human leu	196	15	100.0	9	6	ABJ24784	162P1E6 c
124	15	100.0	9	5	AAE18750	Aae18750	Human leu	197	15	100.0	9	6	ABJ20592	162P1E6 c
125	15	100.0	9	5	AAE18747	Aae18747	Human leu	198	15	100.0	9	6	ABJ21266	162P1E6 c
126	15	100.0	9	5	ABJ12791	Abj12791	Human 125	199	15	100.0	9	6	ABJ21286	162P1E6 c
127	15	100.0	9	5	ABJ12828	Abj12828	Human 125	200	15	100.0	9	6	ABJ22678	162P1E6 c
128	15	100.0	9	5	ABJ13140	Abj13140	Human 125	201	15	100.0	9	6	ABJ23354	162P1E6 c
129	15	100.0	9	5	ABJ12773	Abj12773	Human 125	202	15	100.0	9	6	ABJ20593	162P1E6 c
130	15	100.0	9	5	ABJ13011	Abj13011	Human 125	203	15	100.0	9	6	ABJ22677	162P1E6 c
131	15	100.0	9	5	ABJ12002	Abj12002	Human 125	204	15	100.0	9	6	ABJ23367	162P1E6 c
132	15	100.0	9	5	ABJ12107	Abj12107	Human 125	205	15	100.0	9	6	ABJ21285	162P1E6 c
133	15	100.0	9	5	ABJ12580	Abj12580	Human 125	206	15	100.0	9	6	ABJ21964	162P1E6 c
134	15	100.0	9	5	ABJ12673	Abj12673	Human 125	207	15	100.0	9	6	ABJ21997	162P1E6 c
135	15	100.0	9	5	ABJ12709	Abj12709	Human 125	208	15	100.0	9	6	ABJ22694	162P1E6 c
136	15	100.0	9	5	ABJ13033	Abj13033	Human 125	209	15	100.0	9	6	ABJ24051	162P1E6 c
137	15	100.0	9	5	ABJ13221	Abj13221	Human 125	210	15	100.0	9	6	ABJ24052	162P1E6 c
138	15	100.0	9	5	ABJ12655	Abj12655	Human 125	211	15	100.0	9	6	ABJ21986	162P1E6 c
139	15	100.0	9	5	ABJ13244	Abj13244	Human 125	212	15	100.0	9	6	ABJ22661	162P1E6 c
140	15	100.0	9	5	ABJ12278	Abj12278	Human 125	213	15	100.0	9	6	ABJ22685	162P1E6 c
141	15	100.0	9	5	AAU82561	Aau82561	Llama CDR	214	15	100.0	9	6	ABJ24054	162P1E6 c
142	15	100.0	9	5	AAE17429	Aae17429	Bacteriop	215	15	100.0	9	6	ABJ24063	162P1E6 c
143	15	100.0	9	5	AAU95053	Aau95053	Human nov	216	15	100.0	9	6	ABJ24064	162P1E6 c
144	15	100.0	9	5	AAU79913	Aau79913	Corynebac	217	15	100.0	9	6	ABJ24773	162P1E6 c
145	15	100.0	9	6	ABU73981	Abu73981	Novel pro	218	15	100.0	9	6	ABJ23362	162P1E6 c
146	15	100.0	9	6	ABU77115	Abu77115	Novel pro	219	15	100.0	9	6	ABJ20596	162P1E6 c
147	15	100.0	9	6	ABU72915	Abu72915	Novel pro	220	15	100.0	9	6	ABJ21300	162P1E6 c
148	15	100.0	9	6	ABU73412	Abu73412	Novel pro	221	15	100.0	9	6	ABJ21299	162P1E6 c
149	15	100.0	9	6	ABU73941	Abu73941	Novel pro	222	15	100.0	9	6	ABJ23391	162P1E6 c
150	15	100.0	9	6	ABU75534	Abu75534	Novel pro	223	15	100.0	9	6	ABR16032	Human can
151	15	100.0	9	6	ABU75580	Abu75580	Novel pro	224	15	100.0	9	6	ABR16580	Human can
152	15	100.0	9	6	ABU77167	Abu77167	Novel pro	225	15	100.0	9	6	ABR16053	Human can
153	15	100.0	9	6	ABU77303	Abu77303	Novel pro	226	15	100.0	9	6	ABR16237	Human can
154	15	100.0	9	6	ABU76085	Abu76085	Novel pro	227	15	100.0	9	6	ABR16392	Human can
155	15	100.0	9	6	ABU74451	Abu74451	Novel pro	228	15	100.0	9	6	ABR16767	Human can
156	15	100.0	9	6	ABU76143	Abu76143	Novel pro	229	15	100.0	9	6	ABR17008	Human can
157	15	100.0	9	6	ABU77342	Abu77342	Novel pro	230	15	100.0	9	6	ABR18242	Human can
158	15	100.0	9	6	ABU74459	Abu74459	Novel pro	231	15	100.0	9	6	ABR16662	Human can
159	15	100.0	9	6	ABU75854	Abu75854	Novel pro	232	15	100.0	9	6	ABR17603	Human can
160	15	100.0	9	6	ABU77185	Abu77185	Novel pro	233	15	100.0	9	6	ABR18040	Human can
161	15	100.0	9	6	ABU77284	Abu77284	Novel pro	234	15	100.0	9	6	ABR16574	Human can
162	15	100.0	9	6	ABU73948	Abu73948	Novel pro	235	15	100.0	9	6	ABR17981	Human can
163	15	100.0	9	6	ABU77358	Abu77358	Novel pro	236	15	100.0	9	6	ABR16596	Human can
164	15	100.0	9	6	ABU75552	Abu75552	Novel pro	237	15	100.0	9	6	ABR16791	Human can
165	15	100.0	9	6	ABU75586	Abu75586	Novel pro	238	15	100.0	9	6	ABR16973	Human can
166	15	100.0	9	6	ABU72866	Abu72866	Novel pro	239	15	100.0	9	6	ABR17185	Human can
167	15	100.0	9	6	ABU75058	Abu75058	Novel pro	240	15	100.0	9	6	ABR16369	Human can
168	15	100.0	9	6	ABU77289	Abu77289	Novel pro	241	15	100.0	9	6	ABR16398	Human can
169	15	100.0	9	6	ABU74463	Abu74463	Novel pro	242	15	100.0	9	6	ABR18461	Human can
170	15	100.0	9	6	ABU74506	Abu74506	Novel pro	243	15	100.0	9	6	ABR18644	Human can
171	15	100.0	9	6	ABU76148	Abu76148	Novel pro	244	15	100.0	9	6	ABR16209	Human can

245	15	100.0	9	6	ABR16244	Human Can	AbJ16244	Human Can	318	15	100.0	9	6	ABJ64392	AbJ64392	184P1E2-r
246	15	100.0	9	6	ABJ60991	184P1E2-r	AbJ60991	184P1E2-r	319	15	100.0	9	7	ADC22338	ADC22338	Nuclear 1
247	15	100.0	9	6	ABJ61120	184P1E2-r	AbJ61120	184P1E2-r	320	15	100.0	9	7	ADC53179	ADC53179	Human Cyt
248	15	100.0	9	6	ABJ63446	184P1E2-r	AbJ63446	184P1E2-r	321	15	100.0	9	7	ADD22354	ADD22354	HLA-B46 c
249	15	100.0	9	6	ABJ64742	184P1E2-r	AbJ64742	184P1E2-r	322	15	100.0	9	7	ADD22353	ADD22353	HLA-B46 c
250	15	100.0	9	6	ABJ65479	184P1E2-r	AbJ65479	184P1E2-r	323	15	100.0	10	2	AAK25219	AAK25219	Residues
251	15	100.0	9	6	ABJ61164	184P1E2-r	AbJ61164	184P1E2-r	324	15	100.0	10	2	AAK25224	AAK25224	Residues
252	15	100.0	9	6	ABJ61164	184P1E2-r	AbJ61164	184P1E2-r	325	15	100.0	10	2	AAK28136	AAK28136	Cell-to-c
253	15	100.0	9	6	ABJ61593	184P1E2-r	AbJ61593	184P1E2-r	326	15	100.0	10	2	AAK74985	AAK74985	N-termina
254	15	100.0	9	6	ABJ63538	184P1E2-r	AbJ63538	184P1E2-r	327	15	100.0	10	2	AAW12571	AAW12571	SH2 bindi
255	15	100.0	9	6	ABJ64052	184P1E2-r	AbJ64052	184P1E2-r	328	15	100.0	10	2	AAW12566	AAW12566	SH2 bindi
256	15	100.0	9	6	ABJ60239	184P1E2-r	AbJ60239	184P1E2-r	329	15	100.0	10	2	AAW24171	AAW24171	Peanut al
257	15	100.0	9	6	ABJ61226	184P1E2-r	AbJ61226	184P1E2-r	330	15	100.0	10	2	AAW35338	AAW35338	Rat GDNF
258	15	100.0	9	6	ABJ61227	184P1E2-r	AbJ61227	184P1E2-r	331	15	100.0	10	2	AAK30682	AAK30682	Apo-B100
259	15	100.0	9	6	ABJ61520	184P1E2-r	AbJ61520	184P1E2-r	332	15	100.0	10	2	AAK15253	AAK15253	Peanut al
260	15	100.0	9	6	ABJ65480	184P1E2-r	AbJ65480	184P1E2-r	333	15	100.0	10	2	AAK06987	AAK06987	HLA bindi
261	15	100.0	9	6	ABJ58783	184P1E2-r	AbJ58783	184P1E2-r	334	15	100.0	10	2	AAK07157	AAK07157	HLA bindi
262	15	100.0	9	6	ABJ59836	184P1E2-r	AbJ59836	184P1E2-r	335	15	100.0	10	2	AAW84177	AAW84177	Rat GDNFR
263	15	100.0	9	6	ABJ60280	184P1E2-r	AbJ60280	184P1E2-r	336	15	100.0	10	2	AAK40921	AAK40921	Ara h 1 a
264	15	100.0	9	6	ABJ61119	184P1E2-r	AbJ61119	184P1E2-r	337	15	100.0	10	3	AAK78309	AAK78309	NRD somat
265	15	100.0	9	6	ABJ61731	184P1E2-r	AbJ61731	184P1E2-r	338	15	100.0	10	3	AAK38098	AAK38098	Human ABC
266	15	100.0	9	6	ABJ61805	184P1E2-r	AbJ61805	184P1E2-r	339	15	100.0	10	3	AAK32161	AAK32161	Peptide m
267	15	100.0	9	6	ABJ65478	184P1E2-r	AbJ65478	184P1E2-r	340	15	100.0	10	3	AAK27522	AAK27522	Ara h 1 l
268	15	100.0	9	6	ABJ59488	184P1E2-r	AbJ59488	184P1E2-r	341	15	100.0	10	3	AAK27521	AAK27521	Ara h 1 l
269	15	100.0	9	6	ABJ61594	184P1E2-r	AbJ61594	184P1E2-r	342	15	100.0	10	3	AAK27525	AAK27525	Ara h 1 l
270	15	100.0	9	6	ABJ61662	184P1E2-r	AbJ61662	184P1E2-r	343	15	100.0	10	3	AAK27457	AAK27457	Ara h 1 a
271	15	100.0	9	6	ABJ61933	184P1E2-r	AbJ61933	184P1E2-r	344	15	100.0	10	3	AAK27524	AAK27524	Ara h 1 l
272	15	100.0	9	6	ABJ61936	184P1E2-r	AbJ61936	184P1E2-r	345	15	100.0	10	3	AAK27523	AAK27523	Ara h 1 l
273	15	100.0	9	6	ABJ62332	184P1E2-r	AbJ62332	184P1E2-r	346	15	100.0	10	3	AAK30357	AAK30357	C. elegans
274	15	100.0	9	6	ABJ62688	184P1E2-r	AbJ62688	184P1E2-r	347	15	100.0	10	3	AAK33505	AAK33505	Human imm
275	15	100.0	9	6	ABJ62690	184P1E2-r	AbJ62690	184P1E2-r	348	15	100.0	10	3	AAK23056	AAK23056	Peanut Ar
276	15	100.0	9	6	ABJ63295	184P1E2-r	AbJ63295	184P1E2-r	349	15	100.0	10	4	AAK98322	AAK98322	Human pep
277	15	100.0	9	6	ABJ63699	184P1E2-r	AbJ63699	184P1E2-r	350	15	100.0	10	4	AAK83827	AAK83827	Arabidops
278	15	100.0	9	6	ABJ64741	184P1E2-r	AbJ64741	184P1E2-r	351	15	100.0	10	4	AAK84014	AAK84014	Arabidops
279	15	100.0	9	6	ABJ59837	184P1E2-r	AbJ59837	184P1E2-r	352	15	100.0	10	4	AAK83386	AAK83386	Arabidops
280	15	100.0	9	6	ABJ61941	184P1E2-r	AbJ61941	184P1E2-r	353	15	100.0	10	4	AAK83480	AAK83480	Arabidops
281	15	100.0	9	6	ABJ62687	184P1E2-r	AbJ62687	184P1E2-r	354	15	100.0	10	4	AAK83788	AAK83788	Amino aci
282	15	100.0	9	6	ABJ63821	184P1E2-r	AbJ63821	184P1E2-r	355	15	100.0	10	4	AAK83776	AAK83776	Amino aci
283	15	100.0	9	6	ABJ65557	184P1E2-r	AbJ65557	184P1E2-r	356	15	100.0	10	4	AAK83785	AAK83785	Amino aci
284	15	100.0	9	6	ABJ59879	184P1E2-r	AbJ59879	184P1E2-r	357	15	100.0	10	4	AAU04718	AAU04718	IgE bindi
285	15	100.0	9	6	ABJ50214	184P1E2-r	AbJ50214	184P1E2-r	358	15	100.0	10	4	AAK95854	AAK95854	Human com
286	15	100.0	9	6	ABJ61405	184P1E2-r	AbJ61405	184P1E2-r	359	15	100.0	10	4	AAK95629	AAK95629	Human com
287	15	100.0	9	6	ABJ62308	184P1E2-r	AbJ62308	184P1E2-r	360	15	100.0	10	4	AAK94020	AAK94020	Human com
288	15	100.0	9	6	ABJ62656	184P1E2-r	AbJ62656	184P1E2-r	361	15	100.0	10	4	AAK95615	AAK95615	Human com
289	15	100.0	9	6	ABJ62689	184P1E2-r	AbJ62689	184P1E2-r	362	15	100.0	10	4	AAK94022	AAK94022	Human com
290	15	100.0	9	6	ABJ63165	184P1E2-r	AbJ63165	184P1E2-r	363	15	100.0	10	4	AAK95607	AAK95607	Human com
291	15	100.0	9	6	ABJ64957	184P1E2-r	AbJ64957	184P1E2-r	364	15	100.0	10	4	AAK96948	AAK96948	Human com
292	15	100.0	9	6	ABJ57614	184P1E2-r	AbJ57614	184P1E2-r	365	15	100.0	10	4	AAK95631	AAK95631	Human com
293	15	100.0	9	6	ABJ65556	184P1E2-r	AbJ65556	184P1E2-r	366	15	100.0	10	4	AAK95436	AAK95436	Human com
294	15	100.0	9	6	ABJ59228	184P1E2-r	AbJ59228	184P1E2-r	367	15	100.0	10	4	AAK95617	AAK95617	Human com
295	15	100.0	9	6	ABJ63294	184P1E2-r	AbJ63294	184P1E2-r	368	15	100.0	10	4	AAK13158	AAK13158	Human SCR
296	15	100.0	9	6	ABJ59227	184P1E2-r	AbJ59227	184P1E2-r	369	15	100.0	10	4	AAK85897	AAK85897	Saccharom
297	15	100.0	9	6	ABJ62866	184P1E2-r	AbJ62866	184P1E2-r	370	15	100.0	10	4	AAK86163	AAK86163	Saccharom
298	15	100.0	9	6	ABJ64040	184P1E2-r	AbJ64040	184P1E2-r	371	15	100.0	10	4	AAK86255	AAK86255	Saccharom
299	15	100.0	9	6	ABJ60003	184P1E2-r	AbJ60003	184P1E2-r	372	15	100.0	10	4	AAU05043	AAU05043	Human 19E
300	15	100.0	9	6	ABJ60322	184P1E2-r	AbJ60322	184P1E2-r	373	15	100.0	10	4	ABP23713	ABP23713	HIV A11 m
301	15	100.0	9	6	ABJ63927	184P1E2-r	AbJ63927	184P1E2-r	374	15	100.0	10	4	ABP17504	ABP17504	HIV B27 s
302	15	100.0	9	6	ABJ64669	184P1E2-r	AbJ64669	184P1E2-r	375	15	100.0	10	5	ABE74467	ABE74467	DNA repai
303	15	100.0	9	6	ABJ64740	184P1E2-r	AbJ64740	184P1E2-r	376	15	100.0	10	5	ABE74474	ABE74474	DNA repai
304	15	100.0	9	6	ABJ65177	184P1E2-r	AbJ65177	184P1E2-r	377	15	100.0	10	5	ABU14005	ABU14005	Human 125
305	15	100.0	9	6	ABJ57872	184P1E2-r	AbJ57872	184P1E2-r	378	15	100.0	10	5	ABU13480	ABU13480	Human 125
306	15	100.0	9	6	ABJ59840	184P1E2-r	AbJ59840	184P1E2-r	379	15	100.0	10	5	ABU13523	ABU13523	Human 125
307	15	100.0	9	6	ABJ59229	184P1E2-r	AbJ59229	184P1E2-r	380	15	100.0	10	5	ABU12157	ABU12157	Human 125
308	15	100.0	9	6	ABJ59797	184P1E2-r	AbJ59797	184P1E2-r	381	15	100.0	10	5	ABU11957	ABU11957	Human 125
309	15	100.0	9	6	ABJ62865	184P1E2-r	AbJ62865	184P1E2-r	382	15	100.0	10	5	ABU13600	ABU13600	Human 125
310	15	100.0	9	6	ABJ63131	184P1E2-r	AbJ63131	184P1E2-r	383	15	100.0	10	5	ABU14168	ABU14168	Human 125
311	15	100.0	9	6	ABJ64261	184P1E2-r	AbJ64261	184P1E2-r	384	15	100.0	10	5	ABU11880	ABU11880	Human 125
312	15	100.0	9	6	ABJ60373	184P1E2-r	AbJ60373	184P1E2-r	385	15	100.0	10	5	ABU11694	ABU11694	Human 125
313	15	100.0	9	6	ABJ59878	184P1E2-r	AbJ59878	184P1E2-r	386	15	100.0	10	5	ABU14169	ABU14169	Human 125
314	15	100.0	9	6	ABJ64411	184P1E2-r	AbJ64411	184P1E2-r	387	15	100.0	10	5	ABU12052	ABU12052	Human 125
315	15	100.0	9	6	ABU57870	184P1E2-r	AbJ57870	184P1E2-r	388	15	100.0	10	5	ABU13809	ABU13809	Human 125
316	15	100.0	9	6	ABU60279	184P1E2-r	AbJ60279	184P1E2-r	389	15	100.0	10	5	ABU14170	ABU14170	Human 125
317	15	100.0	9	6	ABU62180	184P1E2-r	AbJ62180	184P1E2-r	390	15	100.0	10	5	ABU13965	ABU13965	Human 125

391	15	100.0	10	5	ABJ13978	Abj13978 Human 125	464	15	100.0	10	6	ABJ20948	Abj20948 162P1E6 C
392	15	100.0	10	5	ABJ05634	Abj05634 Peptide m	465	15	100.0	10	6	ABJ21623	Abj21623 162P1E6 C
393	15	100.0	10	5	AAU78947	AAU78947 CAMP depe	466	15	100.0	10	6	ABJ21638	Abj21638 162P1E6 C
394	15	100.0	10	5	AAU93333	AAU93333 Granulocy	467	15	100.0	10	6	ABJ21641	Abj21641 162P1E6 C
395	15	100.0	10	5	AAU93331	AAU93331 Granulocy	468	15	100.0	10	6	ABJ23023	Abj23023 162P1E6 C
396	15	100.0	10	5	AAU93332	AAU93332 Granulocy	469	15	100.0	10	6	ABJ23716	Abj23716 162P1E6 C
397	15	100.0	10	5	ABG68970	ABG68970 Signature	470	15	100.0	10	6	ABJ22335	Abj22335 162P1E6 C
398	15	100.0	10	5	ABG68958	ABG68958 Signature	471	15	100.0	10	6	ABJ23007	Abj23007 162P1E6 C
399	15	100.0	10	6	ABU73150	Abu73150 Novel pro	472	15	100.0	10	6	ABJ22310	Abj22310 162P1E6 C
400	15	100.0	10	6	ABU73172	Abu73172 Novel pro	473	15	100.0	10	6	ABJ22345	Abj22345 162P1E6 C
401	15	100.0	10	6	ABU75869	Abu75869 Novel pro	474	15	100.0	10	6	ABJ23009	Abj23009 162P1E6 C
402	15	100.0	10	6	ABU74207	Abu74207 Novel pro	475	15	100.0	10	6	ABJ23720	Abj23720 162P1E6 C
403	15	100.0	10	6	ABU75831	Abu75831 Novel pro	476	15	100.0	10	6	ABJ21633	Abj21633 162P1E6 C
404	15	100.0	10	6	ABU76345	Abu76345 Novel pro	477	15	100.0	10	6	ABJ21634	Abj21634 162P1E6 C
405	15	100.0	10	6	ABU73177	Abu73177 Novel pro	478	15	100.0	10	6	ABJ21644	Abj21644 162P1E6 C
406	15	100.0	10	6	ABU74781	Abu74781 Novel pro	479	15	100.0	10	6	ABJ24422	Abj24422 162P1E6 C
407	15	100.0	10	6	ABU78027	Abu78027 Novel pro	480	15	100.0	10	6	ABJ25104	Abj25104 162P1E6 C
408	15	100.0	10	6	ABU73138	Abu73138 Novel pro	481	15	100.0	10	6	ABJ24406	Abj24406 162P1E6 C
409	15	100.0	10	6	ABU74241	Abu74241 Novel pro	482	15	100.0	10	6	ABJ25124	Abj25124 162P1E6 C
410	15	100.0	10	6	ABU74716	Abu74716 Novel pro	483	15	100.0	10	6	ABJ23032	Abj23032 162P1E6 C
411	15	100.0	10	6	ABU75812	Abu75812 Novel pro	484	15	100.0	10	6	ABJ24400	Abj24400 162P1E6 C
412	15	100.0	10	6	ABU73682	Abu73682 Novel pro	485	15	100.0	10	6	ABJ25128	Abj25128 162P1E6 C
413	15	100.0	10	6	ABU74197	Abu74197 Novel pro	486	15	100.0	10	6	ABR16891	Abri16891 Human can
414	15	100.0	10	6	ABU76330	Abu76330 Novel pro	487	15	100.0	10	6	ABR17557	Abri17557 Human can
415	15	100.0	10	6	ABU78114	Abu78114 Novel pro	488	15	100.0	10	6	ABR17883	Abri17883 Human can
416	15	100.0	10	6	ABU75879	Abu75879 Novel pro	489	15	100.0	10	6	ABR18129	Abri18129 Human can
417	15	100.0	10	6	ABU74741	Abu74741 Novel pro	490	15	100.0	10	6	ABR16480	Abri16480 Human can
418	15	100.0	10	6	ABU75855	Abu75855 Novel pro	491	15	100.0	10	6	ABR16095	Abri16095 Human can
419	15	100.0	10	6	ABU76405	Abu76405 Novel pro	492	15	100.0	10	6	ABR18470	Abri18470 Human can
420	15	100.0	10	6	ABU74269	Abu74269 Novel pro	493	15	100.0	10	6	ABR16144	Abri16144 Human can
421	15	100.0	10	6	ABU78166	Abu78166 Novel pro	494	15	100.0	10	6	ABR17078	Abri17078 Human can
422	15	100.0	10	6	ABU74250	Abu74250 Novel pro	495	15	100.0	10	6	ABR16270	Abri16270 Human can
423	15	100.0	10	6	ABU78120	Abu78120 Novel pro	496	15	100.0	10	6	ABR16901	Abri16901 Human can
424	15	100.0	10	6	ABU74721	Abu74721 Novel pro	497	15	100.0	10	6	ABR17124	Abri17124 Human can
425	15	100.0	10	6	ABU74765	Abu74765 Novel pro	498	15	100.0	10	6	ABR17880	Abri17880 Human can
426	15	100.0	10	6	ABU75319	Abu75319 Novel pro	499	15	100.0	10	6	ABR18069	Abri18069 Human can
427	15	100.0	10	6	ABU73196	Abu73196 Novel pro	500	15	100.0	10	6	ABR18289	Abri18289 Human can
428	15	100.0	10	6	ABU75847	Abu75847 Novel pro	501	15	100.0	10	6	ABR16689	Abri16689 Human can
429	15	100.0	10	6	ABU90789	Abu90789 Peptide #	502	15	100.0	10	6	ABR16715	Abri16715 Human can
430	15	100.0	10	6	ABJ23037	Abj23037 162P1E6 C	503	15	100.0	10	6	ABR17308	Abri17308 Human can
431	15	100.0	10	6	ABJ23732	Abj23732 162P1E6 C	504	15	100.0	10	6	ABR17760	Abri17760 Human can
432	15	100.0	10	6	ABJ24426	Abj24426 162P1E6 C	505	15	100.0	10	6	ABR16328	Abri16328 Human can
433	15	100.0	10	6	ABJ24428	Abj24428 162P1E6 C	506	15	100.0	10	6	ABR16523	Abri16523 Human can
434	15	100.0	10	6	ABJ20921	Abj20921 162P1E6 C	507	15	100.0	10	6	ABR16910	Abri16910 Human can
435	15	100.0	10	6	ABJ20949	Abj20949 162P1E6 C	508	15	100.0	10	6	ABR17324	Abri17324 Human can
436	15	100.0	10	6	ABJ20951	Abj20951 162P1E6 C	509	15	100.0	10	6	ABR18328	Abri18328 Human can
437	15	100.0	10	6	ABJ21610	Abj21610 162P1E6 C	510	15	100.0	10	6	ABR18493	Abri18493 Human can
438	15	100.0	10	6	ABJ21636	Abj21636 162P1E6 C	511	15	100.0	10	6	ABR16492	Abri16492 Human can
439	15	100.0	10	6	ABJ24403	Abj24403 162P1E6 C	512	15	100.0	10	6	ABR16545	Abri16545 Human can
440	15	100.0	10	6	ABJ24407	Abj24407 162P1E6 C	513	15	100.0	10	6	ABR16705	Abri16705 Human can
441	15	100.0	10	6	ABJ25102	Abj25102 162P1E6 C	514	15	100.0	10	6	ABR16733	Abri16733 Human can
442	15	100.0	10	6	ABJ25126	Abj25126 162P1E6 C	515	15	100.0	10	6	ABR16764	Abri16764 Human can
443	15	100.0	10	6	ABJ232315	Abj232315 162P1E6 C	516	15	100.0	10	6	ABR18164	Abri18164 Human can
444	15	100.0	10	6	ABJ23006	Abj23006 162P1E6 C	517	15	100.0	10	6	ABR18730	Abri18730 Human can
445	15	100.0	10	6	ABJ23706	Abj23706 162P1E6 C	518	15	100.0	10	6	ABR17145	Abri17145 Human can
446	15	100.0	10	6	ABJ23722	Abj23722 162P1E6 C	519	15	100.0	10	6	ABR16964	Abri16964 Human can
447	15	100.0	10	6	ABJ23723	Abj23723 162P1E6 C	520	15	100.0	10	6	ABR16267	Abri16267 Human can
448	15	100.0	10	6	ABJ20934	Abj20934 162P1E6 C	521	15	100.0	10	6	ABR18094	Abri18094 Human can
449	15	100.0	10	6	ABJ20942	Abj20942 162P1E6 C	522	15	100.0	10	6	ABR17337	Abri17337 Human can
450	15	100.0	10	6	ABJ22329	Abj22329 162P1E6 C	523	15	100.0	10	6	ABR18674	Abri18674 Human can
451	15	100.0	10	6	ABJ23017	Abj23017 162P1E6 C	524	15	100.0	10	6	ABR17102	Abri17102 Human can
452	15	100.0	10	6	ABJ23042	Abj23042 162P1E6 C	525	15	100.0	10	6	ABU52457	Abu52457 Peanut Ar
453	15	100.0	10	6	ABJ20915	Abj20915 162P1E6 C	526	15	100.0	10	6	ABU5454	Abu5454 Peanut Ar
454	15	100.0	10	6	ABJ22313	Abj22313 162P1E6 C	527	15	100.0	10	6	ABU52456	Abu52456 Peanut Ar
455	15	100.0	10	6	ABJ23710	Abj23710 162P1E6 C	528	15	100.0	10	6	ABU52420	Abu52420 Peanut Ar
456	15	100.0	10	6	ABJ25105	Abj25105 162P1E6 C	529	15	100.0	10	6	ABU52453	Abu52453 Peanut Ar
457	15	100.0	10	6	ABJ25121	Abj25121 162P1E6 C	530	15	100.0	10	6	ABU52455	Abu52455 Peanut Ar
458	15	100.0	10	6	ABJ22334	Abj22334 162P1E6 C	531	15	100.0	10	6	ABU69942	Abu69942 Human imm
459	15	100.0	10	6	ABJ25099	Abj25099 162P1E6 C	532	15	100.0	10	6	ABU69786	Abu69786 Human imm
460	15	100.0	10	6	ABJ22325	Abj22325 162P1E6 C	533	15	100.0	10	6	ABJ57989	Abj57989 184P1E2-r
461	15	100.0	10	6	ABJ23727	Abj23727 162P1E6 C	534	15	100.0	10	6	ABJ66010	Abj66010 184P1E2-r
462	15	100.0	10	6	ABJ24421	Abj24421 162P1E6 C	535	15	100.0	10	6	ABJ66088	Abj66088 184P1E2-r
463	15	100.0	10	6	ABJ20918	Abj20918 162P1E6 C	536	15	100.0	10	6	ABJ68260	Abj68260 184P1E2-r

537	15	100.0	10	6	ABJ68383	ABJ68383 184PIE2-r	610	15	100.0	12	2	AAW21260 Hydroxyme
538	15	100.0	10	6	ABJ65782	ABJ65782 184PIE2-r	611	15	100.0	12	2	AAW66777 Cell adhe
539	15	100.0	10	6	ABJ66828	ABJ66828 184PIE2-r	612	15	100.0	12	2	AAW63620 Human HDC
540	15	100.0	10	6	ABJ68261	ABJ68261 184PIE2-r	613	15	100.0	12	2	AAW39461 CD147 bin
541	15	100.0	10	6	ABJ68667	ABJ68667 184PIE2-r	614	15	100.0	12	2	AAW14377 Peptide #
542	15	100.0	10	6	ABJ66681	ABJ66681 184PIE2-r	615	15	100.0	12	2	AAW14378 Peptide #
543	15	100.0	10	6	ABJ66681	ABJ66681 184PIE2-r	616	15	100.0	12	2	AAW15792 Antigenic
544	15	100.0	10	6	ABJ68504	ABJ68504 184PIE2-r	617	15	100.0	12	2	AAW15794 Antigenic
545	15	100.0	10	6	ABJ66532	ABJ66532 184PIE2-r	618	15	100.0	12	2	AAW15794 Antigenic
546	15	100.0	10	6	ABJ68262	ABJ68262 184PIE2-r	619	15	100.0	12	3	AAW38096 Human ABC
547	15	100.0	10	6	ABJ67149	ABJ67149 184PIE2-r	620	15	100.0	12	3	AAW38097 Mouse ABC
548	15	100.0	10	6	ABJ68084	ABJ68084 184PIE2-r	621	15	100.0	12	3	AAW08233 Amino aci
549	15	100.0	10	6	ABJ68170	ABJ68170 184PIE2-r	622	15	100.0	12	4	AAE12454 Dodecamer
550	15	100.0	10	6	ABJ66970	ABJ66970 184PIE2-r	623	15	100.0	12	4	AAE12455 Dodecamer
551	15	100.0	10	6	ABJ67129	ABJ67129 184PIE2-r	624	15	100.0	12	4	AAE07435 Synthetic
552	15	100.0	10	6	ABJ67613	ABJ67613 184PIE2-r	625	15	100.0	12	4	AAE07434 Synthetic
553	15	100.0	10	6	ABJ69398	ABJ69398 184PIE2-r	626	15	100.0	12	5	AAE15769 Synthetic
554	15	100.0	10	6	ABJ66637	ABJ66637 184PIE2-r	627	15	100.0	12	5	AAE15769 Synthetic
555	15	100.0	10	6	ABJ68438	ABJ68438 184PIE2-r	628	15	100.0	12	5	ABW94464 Ubiquitin
556	15	100.0	10	6	ABJ66089	ABJ66089 184PIE2-r	629	15	100.0	12	5	ABW81991 Human iPP
557	15	100.0	10	6	ABJ65781	ABJ65781 184PIE2-r	630	15	100.0	12	5	AAU76778 6A peptid
558	15	100.0	10	6	ABJ66531	ABJ66531 184PIE2-r	631	15	100.0	12	5	AAU76779 6B peptid
559	15	100.0	10	6	ABJ67173	ABJ67173 184PIE2-r	632	15	100.0	12	5	ABW70964 Human col
560	15	100.0	10	6	ABJ67322	ABJ67322 184PIE2-r	633	15	100.0	12	5	AAO22420 Protease
561	15	100.0	10	6	ABJ69062	ABJ69062 184PIE2-r	634	15	100.0	12	6	ABW99618 Peptide d
562	15	100.0	10	6	ABJ69399	ABJ69399 184PIE2-r	635	15	100.0	12	7	ADC44476 Endotheli
563	15	100.0	10	6	ABJ68842	ABJ68842 184PIE2-r	636	15	100.0	12	7	ADC13997 Rheumatol
564	15	100.0	10	6	ABJ69015	ABJ69015 184PIE2-r	637	15	100.0	12	7	ADC13887 Rheumatol
565	15	100.0	10	6	ABJ65964	ABJ65964 184PIE2-r	638	15	100.0	13	2	AAW78694 Human nat
566	15	100.0	10	6	ABJ67567	ABJ67567 184PIE2-r	639	15	100.0	13	2	AAW78694 Human nat
567	15	100.0	10	6	ABJ68623	ABJ68623 184PIE2-r	640	15	100.0	13	2	AAW21047 Human gli
568	15	100.0	10	6	ABJ69166	ABJ69166 184PIE2-r	641	15	100.0	13	2	AAW53679 Enteric n
569	15	100.0	10	6	ABJ66996	ABJ66996 184PIE2-r	642	15	100.0	13	4	AAW80501 PTH2 rece
570	15	100.0	10	6	ABJ67292	ABJ67292 184PIE2-r	643	15	100.0	13	5	ABW79232 Human K+a
571	15	100.0	10	6	ABJ67504	ABJ67504 184PIE2-r	644	15	100.0	13	5	ABW79232 Human K+a
572	15	100.0	10	6	ABJ68169	ABJ68169 184PIE2-r	645	15	100.0	13	5	ABW79250 Human K+a
573	15	100.0	10	6	ABJ68171	ABJ68171 184PIE2-r	646	15	100.0	13	5	ABW79250 Human K+a
574	15	100.0	10	6	ABJ68762	ABJ68762 184PIE2-r	647	15	100.0	13	6	ABW23214 HIV-1 tat
575	15	100.0	10	6	ABJ68841	ABJ68841 184PIE2-r	648	15	100.0	13	6	ABW38758 Human G-p
576	15	100.0	10	6	ABJ69167	ABJ69167 184PIE2-r	649	15	100.0	13	6	ABW38758 Human G-p
577	15	100.0	10	6	ABJ67024	ABJ67024 184PIE2-r	650	15	100.0	13	7	ABW033475 BGS-3 PKC
578	15	100.0	10	6	ABJ69061	ABJ69061 184PIE2-r	651	15	100.0	14	2	ABW033475 BGS-3 PKC
579	15	100.0	10	6	ABW99621	ABW99621 Peptide d	652	15	100.0	14	2	AAW15705 Rev HIV-1
580	15	100.0	10	6	ABW04422	ABW04422 Human HMG	653	15	100.0	14	2	AAW96587 Feline le
581	15	100.0	10	7	ADA07567	ADA07567 Human sec	654	15	100.0	14	2	AAW10314 Murine ga
582	15	100.0	10	7	ADC53180	ADC53180 Human Cyt	655	15	100.0	14	2	AAW59114 FMDV non-
583	15	100.0	10	7	ADD96475	ADD96475 HIV-1 cfo	656	15	100.0	14	2	AAW68346 MHC bindi
584	15	100.0	10	7	ADD96319	ADD96319 HIV-1 cfo	657	15	100.0	14	2	AAW63082 Human imm
585	15	100.0	11	2	AAW28089	AAW28089 Cell-to-c	658	15	100.0	14	2	AAW29728 Feline le
586	15	100.0	11	2	AAW28128	AAW28128 Cell-to-c	659	15	100.0	14	2	AAW87572 Anti-hen
587	15	100.0	11	2	AAW28130	AAW28130 Cell-to-c	660	15	100.0	14	3	AAW68223 Altered M
588	15	100.0	11	2	AAW28132	AAW28132 Cell-to-c	661	15	100.0	14	3	AAW98906 HLA class
589	15	100.0	11	2	AAW12602	AAW12602 SH2 bindi	662	15	100.0	14	3	AAW99015 HLA class
590	15	100.0	11	2	AAW15675	AAW15675 Platelet	663	15	100.0	14	3	AAW52877 Altered M
591	15	100.0	11	3	AAW16616	AAW16616 Phosphoin	664	15	100.0	14	4	AAW96942 Human pep
592	15	100.0	11	3	AAW98542	AAW98542 NCM Igl	665	15	100.0	14	4	AAW97114 Human pep
593	15	100.0	11	3	AAW94681	AAW94681 Human zbi	666	15	100.0	14	4	AAW97000 Human pep
594	15	100.0	11	4	AAW27846	AAW27846 Human p53	667	15	100.0	14	4	AAW96943 Human pep
595	15	100.0	11	4	ABP17507	ABP17507 HIV B27 s	668	15	100.0	14	4	AAW96994 Human pep
596	15	100.0	11	5	ABW74475	ABW74475 DNA repai	669	15	100.0	14	4	AAW98733 Human pep
597	15	100.0	11	5	ABW69342	ABW69342 Human neu	670	15	100.0	14	4	AAW97547 Human pep
598	15	100.0	11	5	ABP54083	ABP54083 Transport	671	15	100.0	14	4	AAW91022 Somatosta
599	15	100.0	11	5	ABP61370	ABP61370 Anti-Chro	672	15	100.0	14	4	AAW00284 Human pro
600	15	100.0	11	5	AAU96240	AAU96240 Class I G	673	15	100.0	14	4	AAW00283 Human pro
601	15	100.0	11	6	AAE34290	AAE34290 Human 5-h	674	15	100.0	14	4	AAW58638 Altered M
602	15	100.0	11	6	AAE34216	AAE34216 Human 5-h	675	15	100.0	14	4	ABW56576 Human SNP
603	15	100.0	11	6	ABW99619	ABW99619 Peptide d	676	15	100.0	14	4	ABW56871 Human SNP
604	15	100.0	11	6	ABW99620	ABW99620 Peptide d	677	15	100.0	14	4	AAW67322 Peptide e
605	15	100.0	11	7	ADC19828	ADC19828 Fluoresce	678	15	100.0	14	4	AAW67304 Peptide e
606	15	100.0	12	1	AAW82895	AAW82895 immunosup	679	15	100.0	14	5	AAW80502 PTH2 rece
607	15	100.0	12	1	AAW82369	AAW82369 immunosup	680	15	100.0	14	5	AAE19420 Human myo
608	15	100.0	12	2	AAW12501	AAW12501 Peptide I	681	15	100.0	14	5	ABJ01158 Human neu
609	15	100.0	12	2	AAW62102	AAW62102 Hydrophil	682	15	100.0	14	6	ABW99617 Peptide d

829	15	100.0	16	6	ABC53747	Abc53747 Novel hum	902	15	100.0	18	4	ABE30933	Abb30933 Peptide #
830	15	100.0	17	1	ARP90644	Apr90644 Signal pe	903	15	100.0	18	4	AM78321	Aam78321 Human bon
831	15	100.0	17	2	AR38622	Ar38622 Sequence	904	15	100.0	18	4	AM69293	Aam69293 Human bon
832	15	100.0	17	2	AR33096	Ar33096 Human cyt	905	15	100.0	18	4	AM65705	Aam65705 Human bra
833	15	100.0	17	2	AR33101	Ar33101 Human cyt	906	15	100.0	18	4	AM56905	Aam56905 Human bra
834	15	100.0	17	2	AR57339	Ar57339 Peptide f	907	15	100.0	18	4	ABG50969	Abg50969 Human liv
835	15	100.0	17	2	AR75822	Ar75822 Antimicro	908	15	100.0	18	4	ABG59929	Abg59929 Human liv
836	15	100.0	17	2	AR67707	Ar67707 HIV-1 Rev	909	15	100.0	18	4	AB80506	Ab80506 PTH2 rece
837	15	100.0	17	2	AR93664	Ar93664 HIV princ	910	15	100.0	18	5	AB74678	Ab74678 Transcript
838	15	100.0	17	2	AM68955	Aam68955 Cytotoxic	911	15	100.0	18	5	ABG47346	Abg47346 Human pep
839	15	100.0	17	2	AX24901	Axy24901 Peptide N	912	15	100.0	18	5	ABG38907	Abg38907 Human pep
840	15	100.0	17	2	AY02812	Aay02812 Fragment	913	15	100.0	18	5	ABG32405	Abg32405 Peptide #
841	15	100.0	17	2	AY36571	Aay36571 Fragment	914	15	100.0	18	5	ABG32403	Abg32403 Peptide #
842	15	100.0	17	2	AY36476	Aay36476 Fragment	915	15	100.0	18	5	ABG32402	Abg32402 Peptide #
843	15	100.0	17	4	AM98330	Aam98330 Human pep	916	15	100.0	18	5	ABG32404	Abg32404 Peptide #
844	15	100.0	17	4	AM67730	Aag67730 Peptide H	917	15	100.0	18	5	ABG71092	Abg71092 Escherich
845	15	100.0	17	4	AM52220	Aam52220 HIV-1 Rev	918	15	100.0	18	5	ABE15901	Aab15901 Mitochond
846	15	100.0	17	4	AM91810	Aab91810 Amyloid b	919	15	100.0	18	5	ABG63642	Abg63642 Human alb
847	15	100.0	17	4	AB91777	Ab91777 Amyloid b	920	15	100.0	18	5	AU78941	Aau78941 Nucleopla
848	15	100.0	17	4	AB43353	Ab43353 Peptide #	921	15	100.0	18	5	AU78938	Aau78938 Thyroid A
849	15	100.0	17	4	AM37207	Aam37207 Peptide #	922	15	100.0	18	5	AU78942	Aau78942 C-fos pep
850	15	100.0	17	4	AM97010	Aau97010 Human C/E	923	15	100.0	18	5	ABG95806	Abg95806 Cell pene
851	15	100.0	17	4	AB11953	Aae11953 Nuclear 1	924	15	100.0	18	6	ARO16674	Aro16674 HIV cell-
852	15	100.0	17	4	AM77086	Aam77086 Human bon	925	15	100.0	18	6	ABP83210	Abp83210 G protein
853	15	100.0	17	4	AM64265	Aam64265 Human bra	926	15	100.0	19	2	ABR61270	Aar61270 19-residu
854	15	100.0	17	4	ABG52037	Abg52037 Human liv	927	15	100.0	19	2	ABR82623	Aar82623 70K autca
855	15	100.0	17	4	AB80505	Ab80505 PTH2 rece	928	15	100.0	19	2	ABW07666	Aaw07666 Bacteriop
856	15	100.0	17	5	AM09993	Aau09993 Human M t	929	15	100.0	19	2	AAW42128	Aaw42128 T-cell ep
857	15	100.0	17	5	ABW74352	Abb74352 Nuclear 1	930	15	100.0	19	2	AAW41434	Aaw41434 Mouse P81
858	15	100.0	17	5	AU78969	Aau78969 Thyroid a	931	15	100.0	19	2	AAW41106	Aay41106 Human can
859	15	100.0	17	5	AU78973	Aau78973 C-fos pep	932	15	100.0	19	2	AAW99306	Aay99306 Human BAI
860	15	100.0	17	5	AG80776	Aag80776 Heptadeca	933	15	100.0	19	3	AY98298	Aay98298 Alpha D p
861	15	100.0	17	5	AB23689	Aae23689 Fluoresce	934	15	100.0	19	3	AY79970	Aay79970 Non-typea
862	15	100.0	17	6	AP59497	Abp59497 Human hep	935	15	100.0	19	3	AY79969	Aay79969 Non-typea
863	15	100.0	17	6	ABJ38987	Abj38987 Linear Ga	936	15	100.0	19	4	AM13713	Aam13713 Peptide #
864	15	100.0	17	6	ABJ38987	Abj38987 Linear Ga	937	15	100.0	19	4	AB32645	Abb32645 Peptide #
865	15	100.0	17	6	ABP82068	Abp82068 G protein	938	15	100.0	19	4	AM26114	Aam26114 Peptide #
866	15	100.0	17	6	ABP82617	Abp82617 G protein	939	15	100.0	19	4	AM84644	Aam84644 Human imm
867	15	100.0	17	6	ABU9879	Abu9879 HIV-1 Rev	940	15	100.0	19	4	AB27493	Abb27493 Human pep
868	15	100.0	17	6	ABU99610	Abb99610 Peptide d	941	15	100.0	19	4	AB18142	Abb18142 Protein #
869	15	100.0	17	6	ABB99609	Abb99609 Peptide d	942	15	100.0	19	4	AM65851	Aam65851 Human bon
870	15	100.0	17	6	ABB82916	Abb82916 HIV-1 rev	943	15	100.0	19	4	AAO04808	Aao04808 Human pol
871	15	100.0	17	6	ADA11740	Ada11740 Human nov	944	15	100.0	19	4	AM53473	Aam53473 Human bra
872	15	100.0	17	6	ADA12021	Ada12021 Human nov	945	15	100.0	19	4	ABG47498	Abg47498 Human liv
873	15	100.0	17	7	ADA07528	Ada07528 Human sec	946	15	100.0	19	4	AM01463	Aam01463 Peptide #
874	15	100.0	17	7	ADC22461	Adc22461 RNA bindi	947	15	100.0	19	4	AB80507	Ab80507 PTH2 rece
875	15	100.0	17	7	ADC22370	Adc22370 Nuclear 1	948	15	100.0	19	5	AU87767	Aau87767 Human epi
876	15	100.0	17	7	ADE11588	Adel1588 HIV-1 Rev	949	15	100.0	19	5	AB74811	Abb74811 Nuclear p
877	15	100.0	17	8	ADJ32095	Adj32095 HIV-1 Rev	950	15	100.0	19	5	ABG35486	Abg35486 Human pep
878	15	100.0	18	1	AP81125	Apr81125 C-fog-rel	951	15	100.0	19	5	ABJ01027	Abj01027 Human bre
879	15	100.0	18	2	AR15706	Ar15706 Rev HIV-1	952	15	100.0	19	5	AU93911	Aau93911 Human P45
880	15	100.0	18	2	AR57845	Ar57845 Vnbeta3-	953	15	100.0	19	5	AE23714	Aae23714 Fluoresce
881	15	100.0	18	2	AR86963	Ar86963 Plant nuc	954	15	100.0	19	5	AE23708	Aae23708 Fluoresce
882	15	100.0	18	2	AM05768	Aaw05768 Presentil	955	15	100.0	19	5	AE23754	Aae23754 Alternati
883	15	100.0	18	2	AAW19796	Aaw19796 Plant nuc	956	15	100.0	19	7	ADC22462	Adc22462 RNA bindi
884	15	100.0	18	2	AM72735	Aaw72735 Nuclear t	957	15	100.0	20	1	AP50414	Aap50414 Swine duo
885	15	100.0	18	2	AM66646	Aaw66646 HSW-2 gly	958	15	100.0	20	1	AP71704	Aap71704 Internal
886	15	100.0	18	2	AY24900	Aay24900 Peptide R	959	15	100.0	20	2	AR10787	Aar10787 S-antigen
887	15	100.0	18	2	AY23683	Aay23683 RNA-bindi	960	15	100.0	20	2	AR10788	Aar10788 S-antigen
888	15	100.0	18	3	AY85064	Aay85064 Immunogen	961	15	100.0	20	2	AR27708	Aar27708 PTH/PTHrp
889	15	100.0	18	3	AY85063	Aay85063 Immunogen	962	15	100.0	20	2	AR33097	Aar33097 Human cyt
890	15	100.0	18	3	AB06400	Ab06400 Randomise	963	15	100.0	20	2	AR33098	Aar33098 Human cyt
891	15	100.0	18	3	AY69731	Aay69731 Labelled-	964	15	100.0	20	2	AR33095	Aar33095 Human cyt
892	15	100.0	18	3	AY92059	Aay92059 CHUK/IKK-	965	15	100.0	20	2	ABG62800	Abg62800 Residues
893	15	100.0	18	3	AY82237	Aay82237 Lambda N	966	15	100.0	20	2	ABG2799	Aar62799 Residues
894	15	100.0	18	3	AY97251	Aay97251 M68 TNFR-	967	15	100.0	20	2	AB47507	Aar47507 Tumour su
895	15	100.0	18	4	AM22114	Aam22114 Peptide #	968	15	100.0	20	2	AB64983	Aar64983 WMLV p15E
896	15	100.0	18	4	AM27366	Aam27366 Novel bon	969	15	100.0	20	2	AB98304	Aar98304 p21WAF1 p
897	15	100.0	18	4	AB03084	Aae03084 Human gen	970	15	100.0	20	2	AB98303	Aar98303 p21WAF1 p
898	15	100.0	18	4	AB36122	Abb36122 Peptide #	971	15	100.0	20	2	AR92279	Aar92279 PTH/PTHrp
899	15	100.0	18	4	AB44520	Abb44520 Peptide #	972	15	100.0	20	2	AAW44227	Aaw44227 Human p21
900	15	100.0	18	4	AM29613	Aam29613 Peptide #	973	15	100.0	20	2	AAW44218	Aaw44218 Human p21
901	15	100.0	18	4	AM38567	Aam38567 Peptide #	974	15	100.0	20	2	AM18245	Aaw18245 PSKH-1 se

975 15 100.0 20 2 AAW42129
 976 15 100.0 20 2 AAW73318
 977 15 100.0 20 2 AAY73183
 978 15 100.0 20 3 AAY78381
 979 15 100.0 20 3 AAY98380
 980 15 100.0 20 3 AAB17271
 981 15 100.0 20 3 AAW90840
 982 15 100.0 20 3 AAW90839
 983 15 100.0 20 3 AAB14194
 984 15 100.0 20 4 AAB80508
 985 15 100.0 20 5 ABE79795
 986 15 100.0 20 5 ABE73349
 987 15 100.0 20 5 ABB81206
 988 15 100.0 20 5 AAE23709
 989 15 100.0 20 5 AAE23715
 990 15 100.0 20 5 ABJ15282
 991 15 100.0 20 6 ABJ38089
 992 15 100.0 20 6 ABP83090
 993 15 100.0 20 6 ABP82101
 994 15 100.0 20 6 ABP83254
 995 15 100.0 20 6 ABP82200
 996 15 100.0 20 6 ABP83185
 997 15 100.0 20 6 ABP82628
 998 15 100.0 20 6 ABP83339
 999 15 100.0 20 7 ABP82828
 1000 15 100.0 20 7 ADC99267

ALIGNMENTS

RESULT 1
 AAW56176
 ID AAW56176 standard; peptide; 3 AA.
 XX
 AC AAW56176;
 XX
 DT 20-JUL-1998 (first entry)
 XX
 DE Anti-inflammatory tripeptide.
 XX
 KW Anti-inflammatory; macrophage inhibitory activity; fibronectin;
 KW T-cell inhibitory activity; adherence; extracellular matrix;
 KW up-regulation; fas receptor expression; inflammation.
 XX
 OS Synthetic.
 XX
 PN WO9809985-A2.
 XX
 PD 12-MAR-1998.
 XX
 PF 03-SEP-1997; 97WO-IL000295.
 XX
 PR 03-SEP-1996; 96US-0025376P.
 PR 20-NOV-1996; 96US-00753141.
 PR 28-MAY-1997; 97US-00864301.
 XX
 PA (YEDA) YEDA RES & DEV CO LTD.
 XX
 PI Eisenbachschwartz M, Beserman P, Hirschberg DL;
 XX
 DR WPI; 1998-193550/17.
 XX
 PT Anti-inflammatory peptides and derivatives - used for treating, e.g.
 PT arthritis, ulcerative colitis, auto-immune disease, allergy asthma,
 PT shock, HIV infection, transplant rejection or Alzheimer's disease.
 XX
 PS Claim 3; Page 34; 42pp; English.
 CC
 CC AAW56171-248 represent anti-inflammatory tripeptides of the invention.
 CC They are derived from the formulae: Xaa-Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-
 CC Glu, or Glu-arg-Xaa, where Xaa = any amino acid residue. Cyclic
 CC derivatives of the peptides also function as anti-inflammatory agents.

CC The peptides can be covalently linked to one another either directly or
 CC through a spacer. The peptides and their derivatives have macrophage
 CC inhibitory and T-cell inhibitory activity and thus, anti-inflammatory
 CC activity. The peptides and compositions have anti-immune activity, i.e.
 CC inhibitory effects against a cellular and humoral immune response,
 CC including a response not associated with inflammation. The peptides also
 CC inhibit the ability of macrophages and T-cells to adhere to extracellular
 CC matrix components and fibronectin, as well as up-regulated fas receptor
 CC expression in T-cells. They can be used to inhibit unwanted immune
 CC reaction and inflammation
 XX
 SQ Sequence 3 AA;
 Query Match 100.0%; Score 15; DB 2; Length 3;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 RER 3
 DB 1 RER 3
 RESULT 2
 AAW48192
 ID AAW48192 standard; peptide; 4 AA.
 XX
 AC AAW48192;
 XX
 DT 30-JUN-1998 (first entry)
 XX
 DE Conantokin peptide derivative.
 XX
 KW Conantokin; predatory cone snail; treatment; neurologic disorder;
 KW psychiatric disorder; anticonvulsant; neuroprotective;
 KW analgesic. HIV infection; ophthalmic indication; memory; learning defect;
 KW cognitive defect.
 XX
 OS Synthetic.
 XX
 FH Key Location/Qualifiers
 FT Modified-site 4 /note= "gamma-carboxyglutamic acid"
 XX
 PN WO9803541-A1.
 XX
 PD 29-JAN-1998.
 XX
 PF 21-JUL-1997; 97WO-US012618.
 XX
 PR 22-JUL-1996; 96US-00684742.
 XX
 PA (UTAH) UNIV UTAH RES FOUND.
 PA (COGN-) COGNEX INC.
 XX
 PI Abogadie FC, Cruz LJ, Olivera BM, Walker C, Colledge C;
 PI Hilliard DR, Jimenez E, Laver RT, Zhou L, Shen GS, McCabe RT;
 PI Rivier JE;
 XX
 DR WPI; 1998-120694/11.
 XX
 PT New conantokin peptide(s) - useful for e.g. treating neurologic or
 PT psychiatric disorders, or the management of pain.
 XX
 PS Claim 15; Page 98; 122pp; English.
 CC
 CC The present sequence is a conantokin peptide derivative, which can be
 CC used to treat neurologic and psychiatric disorders, e.g. as an
 CC anticonvulsant, neuroprotective or analgesic agent. Neurologic and
 CC psychiatric disorders include epilepsy, convulsions, neurotoxic injury
 CC (associated with conditions of hypoxia, anoxia or ischaemia, which
 CC typically follow stroke, cerebrovascular accident, brain or spinal cord
 CC trauma, myocardial infarct, physical trauma, drowning, suffocation,
 CC perinatal asphyxia or hypoglycaemic events), neurodegeneration

CC (associated with Alzheimer's disease, senile dementia, Amyotrophic
 CC Lateral Sclerosis, Multiple Sclerosis, Parkinson's disease, Huntington's
 CC disease, Down's Syndrome, Korsakoff's disease, schizophrenia, AIDS
 CC dementia, multi-infarct dementia, Binswanger dementia and neuronal damage
 CC associated with uncontrolled seizures), chemical toxicity (such as
 CC addition, and morphine, opiate, opioid and barbiturate tolerance), pain
 CC (acute, chronic, migraine), anxiety, major depression, manic-depressive
 CC illness, obsessive-compulsive disorder, schizophrenia and mood disorders
 CC (such as bipolar disorder, unipolar depression, dysthymia and seasonal
 CC affective disorder) and dystonia (movement disorder), sleep disorder,
 CC muscle relaxation and urinary incontinence. The peptide can also be used
 CC to treat HIV infection, ophthalmic indication and memory, learning or
 CC cognitive defects
 XX
 SQ Sequence 4 AA;

Query Match 100.0%; Score 15; DB 2; Length 4;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 1 RER 3

RESULT 3
 ID AAW49974 standard; peptide; 4 AA.
 XX AAW49974;
 AC AAW49974;
 DT 30-JUN-1998 (first entry)
 XX
 DE Conantokin peptide derivative.
 DE
 KW Conantokin; predatory cone snail; treatment; neurologic disorder;
 KW psychiatric disorder; anticonvulsant; neuroprotective;
 KW analgesic. HIV infection; ophthalmic indication; memory; learning defect;
 KW cognitive defect.
 XX
 OS Synthetic.
 PH Key Location/Qualifiers
 FT Modified-site 4 /note= "gamma-carboxyglutamic acid"
 FT
 XX WO9803189-A1.
 PN
 XX 29-JAN-1998.
 PD
 XX 21-JUL-1997; 97WO-US012652.
 PF
 XX 22-JUL-1996; 96US-00684750.
 PR 06-DEC-1996; 96US-00762377.
 XX
 PA (COGN-) COGNETIX INC.
 XX
 XX McCabe RT, Zhou L, Layer RT;
 XX WPI; 1998-120469/11.
 XX
 XX Use of conantokin peptide(s) - for treating disorders involving excessive
 PT excitation of nerve cells by excitatory amino acids or agonists of the N-
 PT methyl-D-aspartate receptor.
 XX
 PS Example 19; Page 73; 122pp; English.
 XX
 XX The present sequence is a conantokin peptide derivative, which can be
 CC used to treat neurologic and psychiatric disorders, e.g. as an
 CC anticonvulsant, neuroprotective or analgesic agent. Neurologic and
 CC psychiatric disorders include epilepsy, convulsions, neurotoxic injury
 CC (associated with conditions of hypoxia, anoxia or ischaemia, which
 CC typically follow stroke, cerebrovascular accident, brain or spinal cord

CC trauma, myocardial infarct, physical trauma, drowning, suffocation,
 CC perinatal asphyxia or hypoglycaemic events), neurodegeneration
 CC (associated with Alzheimer's disease, senile dementia, Amyotrophic
 CC Lateral Sclerosis, Multiple Sclerosis, Parkinson's disease, Huntington's
 CC disease, Down's Syndrome, Korsakoff's disease, schizophrenia, AIDS
 CC dementia, multi-infarct dementia, Binswanger dementia and neuronal damage
 CC associated with uncontrolled seizures), chemical toxicity (such as
 CC addition, and morphine, opiate, opioid and barbiturate tolerance), pain
 CC (acute, chronic, migraine), anxiety, major depression, manic-depressive
 CC illness, obsessive-compulsive disorder, schizophrenia and mood disorders
 CC (such as bipolar disorder, unipolar depression, dysthymia and seasonal
 CC affective disorder) and dystonia (movement disorder), sleep disorder,
 CC muscle relaxation and urinary incontinence. The peptide can also be used
 CC to treat HIV infection, ophthalmic indication and memory, learning or
 CC cognitive defects
 XX
 SQ Sequence 4 AA;

Query Match 100.0%; Score 15; DB 2; Length 4;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 1 RER 3

RESULT 4
 ID AAB24196 standard; peptide; 4 AA.
 XX AAB24196;
 AC AAB24196;
 DT 02-FEB-2001 (first entry)
 XX
 DE Dual peptide amino acid sequence SEQ ID NO:1.
 DE
 KW Dual peptide; antithematic; antiarthritic; antiallergic; antiimmune;
 KW antiinflammatory; antiasthmatic; dermatological; immunosuppressive;
 KW antiarteriosclerotic; cardiant; antitumor; hepatotropic; vulnary;
 KW ophthalmological; antiparkinsonian; nootropic; neuroprotective;
 KW cerebroprotective; gynaecological; anticonvulsant; tranquilliser;
 KW antibacterial; cytostatic; inflammation; rheumatoid arthritis;
 KW graft rejection; transplantation; macrophage migration; immune response.
 XX
 OS Synthetic.
 XX US6126939-A.
 PN
 XX 03-OCT-2000.
 PD
 XX 28-MAY-1997; 97US-00864301.
 PF
 XX 03-SEP-1996; 96US-0025376P.
 PR 20-NOV-1996; 96US-0031191P.
 XX 20-NOV-1996; 96US-00753141.
 XX
 XX (YEDA) YEDA RES & DEV CO LTD.
 XX
 XX Eisenbach-Schwartz M, Hirschberg DL, Beerman P;
 XX WPI; 1998-193550/17.
 XX
 XX Anti-inflammatory peptides and derivatives - used for treating, e.g.
 PT arthritis, ulcerative colitis, auto-immune disease, allergy asthma,
 PT shock, HIV infection, transplant rejection or Alzheimer's disease.
 XX
 XX Disclosure; Col 9; 17pp; English.
 PS
 XX The present invention describes a pure anti-inflammatory dipeptide (I)
 CC comprising the sequence of Glu-Arg. (I) can have antirheumatic,
 CC antiarthritic, antiallergic, antiinflammatory, antiimmune, antiasthmatic,
 CC dermatological, immunosuppressive, antiarteriosclerotic, cardiant,

CC antiulcer, hepatotropic, ophthalmological, antiparkinsonian, vulnerary,
 CC neurotropic, neuroprotective, cerebroprotective, gynaecological,
 CC anticonvulsant, tranquiliser, antibacterial and cytostatic activities.
 CC (I) can be used as inhibitors of macrophage migration and/or macrophage
 CC phagocytic activity and inflammation in animals, preferably mammals,
 CC including human. It is used as inhibitors of T cell adhesive activity in
 CC mammals and for the inhibition of an immune response not associated with
 CC inflammation. It is also used for restoration of immune privilege at
 CC immune privileged sites and in the treatment of or amelioration of
 CC inflammatory symptoms in any disease, condition or disorder, e.g.
 CC rheumatoid arthritis. It also prevents and/or treats graft rejection in
 CC cases of transplantation of natural or artificial cells, tissue, and
 CC organs, e.g. cornea, bone marrow, organs, lenses, pacemakers, natural and
 CC artificial skin tissue. (II) inhibits the macrophage activity and has
 CC macrophage migration and/or macrophage phagocytic inhibitory activity as
 CC assessed in vitro assay. It also inhibits T cells and has T cell
 CC inhibitory activity. The present sequence represents a dual peptide which
 CC is used in the exemplification of the present invention

XX
 SQ Sequence 4 AA;

Query Match 100.0%; Score 15; DB 2; Length 4;
 Best Local Similarity 100.0%; Pred. No. 1.4e-06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 2 RER 4

RESULT 5
 AAY71269
 ID AAY71269 standard; peptide; 4 AA.

AC AAY71269;

DT 21-SEP-2000 (first entry)

DE Bovine chondromodulin (Chm)-I protein processing signal sequence, RERR.

XX Chondromodulin-like protein; Zchm1; human; chromosome 11p15.4; cancer;
 KW cell differentiation regulator; osteoblast proliferation stimulator;
 KW cytostatic; diagnostic; therapeutic; polypeptide-toxin fusion protein;
 KW class II cell surface protein; transmembrane domain; gene therapy;
 KW targeted cell inhibition; bovine; chondromodulin-I; Chm-I.

XX Bos sp.

OS WO200029579-A1.

PN 25-MAY-2000.

PF 12-NOV-1999; 99WO-US026909.

PR 13-NOV-1998; 98US-00191986.

PA (ZYMO) ZYMOGENETICS INC.

XX Lok S, Presnell SR;

PI WPI; 2000-387792/33.

XX Polynucleotide encoding mammalian chondromodulin-like polypeptide useful
 PT for gene therapy of various disorders by regulating growth or
 PT differentiation of cells especially cancer cells.

XX Disclosure; Page 1; 87pp; English.

CC The present sequence is the bovine chondromodulin-I (Chm-I) protein
 CC processing signal sequence RERR, that precedes the mature protein
 CC sequence. Bovine Chm-I has sequence homology to human chondromodulin-
 CC like protein, Zchm-1. The Zchm1 locus is mapped to chromosome 11p15.4. It
 CC functions as a cell differentiation regulator and osteoblast

CC proliferation stimulator. Zchm1 can be used as growth or differentiation
 CC regulator for cells, especially mesenchymal, myogenic, chondrogenic or
 CC endothelial cells. Zchm1 proteins or antibodies are useful for
 CC identifying or treating tissues or organs expressing the anti-
 CC complementary molecule, e.g., receptor or antigen. The Zchm1 polypeptides
 CC conjugated to drugs, radionuclides and toxins are useful for in vivo
 CC diagnostic or therapeutic applications and polypeptide-toxin fusion
 CC proteins are useful for targeted cell or tissue inhibition or ablation
 CC for treating various disorders, especially cancer. It is useful for gene
 CC therapy of disorders associated with altered Zchm1 activity

XX

SQ Sequence 4 AA;
 Query Match 100.0%; Score 15; DB 3; Length 4;
 Best Local Similarity 100.0%; Pred. No. 1.4e-06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 1 RER 3

RESULT 6
 AAG79029
 ID AAG79029 standard; peptide; 4 AA.

XX AAG79029;

XX 10-DEC-2001 (first entry)

DE Amino acid sequence of conantokin S1 domain III.

XX Conantokin; cone snail; nerve cell excitation; NMDA receptor; epilepsy;
 KW N-methyl-D-aspartate receptor; pain; psychiatric disorder;
 KW neurotoxic injury; hypoxia; anoxia; ischemia; neurodegeneration;
 KW chemical toxicity; addiction; drug craving; psychiatric disorder;
 KW anxiety; depression; obsessive compulsive disorder; schizophrenia;
 KW mood disorder; ophthalmic disorder; neurological disorder; dystonia;
 KW sleep disorder; muscle relaxation; urinary incontinence;
 KW cognition enhancement; HIV infection.

XX Conus sulcatus.

XX Key Location/Qualifiers

FT Modified-site 4 /note= "gamma-carboxyglutamic acid"

XX US6277825-B1.

XX 21-AUG-2001.

XX 20-JUL-1999; 99US-00357141.

XX 22-JUL-1996; 96US-00684750.

XX 06-DEC-1996; 96US-00762377.

XX 21-JUL-1997; 97WO-US012652.

XX 10-FEB-1999; 99US-00142076.

XX 01-APR-1999; 99US-00283277.

XX (UTAH) UNIV UTAH RES FOUND.

XX (COGN-) COGNETIX INC.

XX Olivera BM, McIntosh JM, McCabe RT, Layer RT, Zhou L;

XX WPI; 2001-601377/68.

XX Use of conantokin peptide or its derivatives or a conantokin peptide
 PT chimera for treating disorders e.g. migraine.

XX Claim 9; Col 80; 60pp; English.

XX AAG79012-43 and AAG790054-56 represent domains of conantokin peptides.
 CC Conantokin differ from conotoxins, in that they contain gamma-

CC carboxylglutamic acid. The conantokins are derived from the venom of cone
 CC snails. They are used for the treatment of disorders in which the
 CC pathophysiology involves excessive excitation of nerve cells by
 CC excitatory amino acids or agonist of N-methyl-D-aspartate (NMDA)
 CC receptor. The conantokin peptides are used for the treatment of disorders
 CC such as pain; neurologic or psychiatric disorders such as epilepsy; for
 CC reducing neurotoxic injury associated with conditions of hypoxia, anoxia
 CC or ischemia; for treating neurodegeneration; for treating chemical,
 CC toxicity such as addiction, drug craving, alcohol abuse, morphine, opioid
 CC and barbiturate tolerance; for treating psychiatric disorders such as
 CC anxiety, major depression, manic-depression illness, obsessive compulsive
 CC disorder, schizophrenia or mood disorder; for treating ophthalmic
 CC disorder; for treating additional neurological disorders e.g. dystonia,
 CC sleep disorder, muscle relaxation and urinary incontinence; for
 CC memory/cognition enhancement; for treating HIV infection
 XX
 SQ Sequence 4 AA;

Query Match 100.0%; Score 15; DB 4; Length 4;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
 ||||
 Db 1 RER 3

RESULT 7
 AAEE16624
 ID AAEE16624 standard; peptide; 4 AA.
 AC AAEE16624;
 XX
 XX 09-APR-2002 (first entry)
 XX
 XX Peptide of human KCNQ5 S4 membrane-spanning domain.
 DE
 XX Human; potassium channel polypeptide; KCNQ5; pain; migraine; stroke;
 KW dementia; trauma; epilepsy; seizure; amyotrophic lateral sclerosis; ALS;
 KW multiple sclerosis; MS; Parkinson's disease; ataxia; depression;
 KW anxiety disorder; bipolar disorder; sleep disorder; eating disorder;
 KW addiction; myokymia; Alzheimer's disease; age-associated memory loss;
 KW learning deficiency; cognitive disorder; motor disease; neuron disease;
 KW neurophysiological disorder; neuropsychological disorder; asthma;
 KW neuron cell death; brain tumour; gene therapy; antisense therapy;
 KW synaptic transmission; S4 membrane-spanning domain;
 KW electrical excitability.
 XX
 OS Homo sapiens.
 XX
 XX WO200192526-A1.
 PN
 XX 06-DEC-2001.
 PD
 XX
 XX 24-MAY-2001; 2001WO-US017314.
 PF
 XX
 XX 26-MAY-2000; 2000US-0207389P.
 PR
 XX (BRIM) BRISTOL-MYERS SQUIBB CO.
 PA
 XX Dworetzky SI, Ramanathan CS, Trojnecki JT, Boissard OG;
 PI Grikboff VK;
 XX
 XX WPI; 2002-122069/16.
 DR
 XX Novel potassium channel polypeptide, KCNQ5 and polynucleotide encoding
 PT it, for diagnosing, treating and identifying modulators useful in
 PT treating neurological, neuropsychological and neuropsychological
 PT diseases.
 XX
 XX Disclosure; Page 21; 128pp; English.
 PS
 XX The invention relates to potassium channel polypeptides referred to as

CC KCNQ5 and nucleic acid molecules encoding such polypeptides. KCNQ5
 CC polypeptides are useful for identifying compounds that modulate their
 CC biological activity. The compounds identified and KCNQ5 polynucleotides
 CC are useful for treating acute and chronic pain, migraine, acute stroke,
 CC dementia, trauma, epilepsy, seizure, amyotrophic lateral sclerosis (ALS),
 CC multiple sclerosis (MS), Parkinson's disease, ataxia, anxiety disorders,
 CC depression, bipolar disorders, sleep disorders, eating disorders,
 CC addiction, myokymia, Alzheimer's disease, age-associated memory loss,
 CC learning deficiencies, cognitive disorders and motor neuron diseases. The
 CC nucleic acid molecules of the invention are further useful for treating
 CC neurophysiological, neuropsychological disorders, asthma, neuron cell
 CC death and brain tumours. They are also used in gene therapy and antisense
 CC therapy. KCNQ5 polypeptides modulate synaptic transmission and electrical
 CC excitability in the brain and are useful for generating antibodies. They
 CC are also useful to affinity purify biological effectors from biological
 CC materials e.g. disease tissues or cells. The present sequence is peptide
 CC of human KCNQ5 S4 membrane-spanning domain
 XX
 SQ Sequence 4 AA;

Query Match 100.0%; Score 15; DB 5; Length 4;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
 ||||
 Db 2 RER 4

RESULT 8
 ABB99613
 ID ABB99613 standard; peptide; 4 AA.
 AC ABB99613;
 XX
 XX 28-MAR-2003 (first entry)
 DT
 XX Peptide derived from human amyloid precursor protein (APP).
 DE
 XX Amyloid precursor protein; APP; protein derivative;
 KW neurodegenerative disease; Alzheimer's disease; cognitive enhancer.
 KW
 OS Synthetic.
 OS Homo sapiens.
 XX WO200283729-A2.
 PN
 XX 24-OCT-2002.
 PD
 XX 17-APR-2002; 2002WO-GB001769.
 PF
 XX 18-APR-2001; 2001GB-00009558.
 PR 17-AUG-2001; 2001GB-00020084.
 PR 30-NOV-2001; 2001US-00998491.
 PR 28-MAR-2002; 2002GB-00007387.
 XX (UYOP-) UNIV OPEN.
 PA
 XX Mileusnic R, Rose SPR;
 PI WPI; 2003-111814/10.
 DR
 XX Derivatives of polypeptides, useful for treating neurodegenerative
 PT disease e.g. Alzheimer's disease, comprises one functional amino acid
 PT residue or derivative protected by a protective group.
 PT
 XX Claim 74; Page 65; 85pp; English.
 PS
 XX The present sequence is derived from amyloid precursor protein (APP).
 CC Derivatives of the invention are based on APP sequences. The
 CC specification describes a derivative of a polypeptide in which at least
 CC one functional group of at least one amino acid residue or derivative is
 CC protected by a protective group. This derivative is of the formula given

CC in ABB99625. The derivative is useful in medicine and in the preparation
 CC of a medicament for use in the treatment of a neurodegenerative disease
 CC e.g. Alzheimer's disease. It is also useful as a cognitive enhancer

XX SQ Sequence 4 AA;

Query Match 100.0%; Score 15; DB 6; Length 4;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 RER 3
 DB 1 RER 3

RESULT 9

ID ABB99614 standard; peptide; 4 AA.

XX AC ABB99614;

XX DT 28-MAR-2003 (first entry)

XX DE Peptide derived from human amyloid precursor protein (APP).

XX KW Amyloid precursor protein; APP; protein derivative;

XX KW neurodegenerative disease; Alzheimer's disease; cognitive enhancer.

XX OS Synthetic.

XX OS Homo sapiens.

XX FN W0200283729-A2.

XX PD 24-OCT-2002.

XX PF 17-APR-2002; 2002WO-GH001769.

XX PR 18-APR-2001; 2001GB-00009558.

XX PR 17-AUG-2001; 2001GB-00020084.

XX PR 30-NOV-2001; 2001US-00998491.

XX PR 28-MAR-2002; 2002GB-00007387.

XX PA (UYOP-) UNIV OPEN.

XX PI Mileusnic R, Rose SPR;

XX DR WPI; 2003-111814/10.

XX PT Derivatives of polypeptides, useful for treating neurodegenerative
 PT disease e.g. Alzheimer's disease, comprises one functional amino acid
 PT residue or derivative protected by a protective group.

XX PS Disclosure; Page 1; 85pp; English.

XX CC The present sequence is derived from amyloid precursor protein (APP).
 CC Derivatives of the invention are based on APP sequences. The
 CC specification describes a derivative of a polypeptide in which at least
 CC one functional group of at least one amino acid residue or derivative is
 CC protected by a protective group. This derivative is of the formula given
 CC in ABB99625. The derivative is useful in medicine and in the preparation
 CC of a medicament for use in the treatment of a neurodegenerative disease
 CC e.g. Alzheimer's disease. It is also useful as a cognitive enhancer

XX SQ Sequence 4 AA;

Query Match 100.0%; Score 15; DB 6; Length 4;

Best Local Similarity 100.0%; Pred. No. 1.4e+06;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 RER 3

DB 2 RER 4

RESULT 10

ID AAR62114 standard; peptide; 5 AA.

XX AC AAR62114;

XX DT 25-MAR-2003 (revised)

XX DT 27-APR-1995 (first entry)

XX DE Hydrophilic motif from U1 snRNP 70K protein.

XX KW Small ribonucleoprotein complex; U1 snRNP; 70K protein; epitope;
 KW autoantibody; immunoinfective cluster virus; nuclear protein antigen;
 KW systemic rheumatic disorder; human immunodeficiency virus; HIV-1.

XX OS Homo sapiens.

XX PN W09420141-A1.

XX PD 15-SEP-1994.

XX PF 10-MAR-1994; 94WO-US002631.

XX PR 11-MAR-1993; 93US-00029850.

XX PA (UYSC-) UNIV SOUTHERN CALIFORNIA.

XX PI Douvas A, Takehana Y, Ehresmann G;

XX DR WPI; 1994-302689/37.

XX PT Methods for treating immunoinfective cluster virus infections - utilise
 PT antibodies or fragments characteristic of auto antibodies produced by
 PT patients with rheumatic disorders.

XX PS Disclosure; Page 8; 106pp; English.

XX CC The sequence RERRR (AAR62113) is a preferred example of an alternating
 CC acidic/basic amino acid, hydrophilic epitope motif, found in the U1 snRNP
 CC 70K protein. It also occurs as RRRER and RRRER (AAR62114 and AAR62115)
 CC in the 70K protein. The motif is also found in similar form in
 CC immunoinfective cluster viruses. The motif serves as an epitope for anti-
 CC viral antibodies and also for autoantibodies which occur in high titre in
 CC patients suffering from systemic rheumatic disorders. Sera from such
 CC patients could be used for treatment of immunoinfective cluster virus
 CC (e.g. HIV, EBV, rubella virus) infections. (Updated on 25-MAR-2003 to
 CC correct PN field.)

XX SQ Sequence 5 AA;

Query Match 100.0%; Score 15; DB 2; Length 5;

Best Local Similarity 100.0%; Pred. No. 1.4e+06;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 RER 3

DB 2 RER 4

RESULT 11

ID AAR62113 standard; peptide; 5 AA.

XX AC AAR62113;

XX DT 25-MAR-2003 (revised)

XX DT 27-APR-1995 (first entry)

XX DE Hydrophilic motif from U1 snRNP 70K protein.

XX KW Small ribonucleoprotein complex; U1 snRNP; 70K protein; epitope;
 KW autoantibody; immunoinfective cluster virus; nuclear protein antigen;

KW systemic rheumatic disorder; human immunodeficiency virus; HIV-1.
 XX Homo sapiens.
 XX WO9420141-A1.
 XX PD 15-SEP-1994.
 XX PF 10-MAR-1994; 94WO-US002631.
 XX PR 11-MAR-1993; 93US-00029850.
 XX PA (UYSC-) UNIV SOUTHERN CALIFORNIA.
 XX PI Douvas A, Takehana Y, Ehresmann G;
 XX WPI; 1994-302689/37.
 XX Methods for treating immunoinfective cluster virus infections - utilise
 PT antibodies or fragments characteristic of auto antibodies produced by
 PT patients with rheumatic disorders.
 XX Claim 13; Page 78; 106pp; English.
 XX The sequence RERRR (AAR62113) is a preferred example of an alternating
 CC acidic/basic amino acid, hydrophilic epitope motif, found in the UI snRNP
 CC 70K protein. It also occurs as RRERE and EREER (AAR62114 and AAR62115)
 CC in the 70K protein. The motif is also found in similar form in
 CC immunoinfective cluster viruses. The motif serves as an epitope for anti-
 CC viral antibodies and also for autoantibodies which occur in high titre in
 CC patients suffering from systemic rheumatic disorders. Sera from such
 CC patients could be used for treatment of immunoinfective cluster virus
 CC (e.g. HIV, EBV, rubella virus) infections. (Updated on 25-MAR-2003 to
 CC correct PN field.)
 XX Sequence 5 AA;
 SQ Query Match 100.0%; Score 15; DB 2; Length 5;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 OY 1 RER 3
 DB 1 RER 3
 RESULT 12
 AAR62154
 ID AAR62154 standard; peptide; 5 AA.
 AC AAR62154;
 XX 27-AUG-2003 (revised)
 DT 25-MAR-2003 (revised)
 DT 02-MAY-1995 (first entry)
 XX Basic/acidic motif from HIV-1 gp120/41 and UI snRNP 70K protein.
 XX Small ribonucleoprotein complex; UI snRNP; 70K protein; epitope;
 KW autoantibody; immunoinfective cluster virus; nuclear protein antigen;
 KW systemic rheumatic disorder; human immunodeficiency virus; HIV-1;
 KW systemic lupus erythematosus; mixed connective tissue disease;
 KW scleroderma; glycoprotein 120; glycoprotein 41.
 XX Homo sapiens.
 OS Human immunodeficiency virus 1.
 XX WO9420141-A1.
 XX PD 15-SEP-1994.
 XX PF 10-MAR-1994; 94WO-US002631.
 XX

PR 11-MAR-1993; 93US-00029850.
 XX (UYSC-) UNIV SOUTHERN CALIFORNIA.
 XX PI Douvas A, Takehana Y, Ehresmann G;
 XX WPI; 1994-302689/37.
 XX Methods for treating immunoinfective cluster virus infections - utilise
 PT antibodies or fragments characteristic of auto antibodies produced by
 PT patients with rheumatic disorders.
 XX Disclosure; Page 59; 106pp; English.
 XX The hydrophilic C-terminal regions of UI snRNP 70K protein and HIV-1 gp41
 CC share extensive homologies. These include the repeating RDRR (AAR62153)
 CC motif and a block of alternating basic and acidic residues beginning at
 CC positions 513 and 732 of 70K and gp41, respectively. In this block, 11 of
 CC 18 of the 70K amino acids are identical to gp41, and 3 more represent
 CC conservative substitutions of Glu and Asp. Configurations of alternating
 CC basic and acidic amino acids (AAR62153- AAR62156) are antigenic to anti-
 CC UI antibodies. Such autoantibodies occur in the systemic rheumatoid
 CC disorders of mixed connective tissue disease, scleroderma and systemic
 CC lupus erythematosus and can be used to neutralise HIV-1. (Updated on 25-
 CC MAR-2003 to correct PN field.) (Updated on 27-AUG-2003 to correct OS
 CC field.)
 XX Sequence 5 AA;
 SQ Query Match 100.0%; Score 15; DB 2; Length 5;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 OY 1 RER 3
 DB 1 RER 3
 RESULT 13
 AAR54661
 ID AAR54661 standard; peptide; 5 AA.
 AC AAR54661;
 XX 25-MAR-2003 (revised)
 DT 29-NOV-1994 (first entry)
 XX Native secreted amyloid precursor protein (APP) peptide.
 DE Amyloid precursor protein; Alzheimer's disease; neuron growth.
 XX Synthetic.
 OS WO9409808-A1.
 XX 11-MAY-1994.
 XX 23-OCT-1992; 92WO-US009070.
 XX 23-OCT-1992; 92WO-US009070.
 XX (REGC) UNIV CALIFORNIA.
 XX Saitoh T;
 XX WPI; 1994-167118/20.
 XX Peptide(s) and analogues based on amyloid precursor protein - used for
 PT promoting neuronal growth in conditions involving damage to neurons or in
 PT treating Alzheimer's Disease etc.
 XX Claim 1; Page 5; 116pp; English.
 XX

CC This sequence corresponds to AA 328-332 of amyloid precursor protein.
 CC This peptide, which is smaller than a native APP, retains at least some
 CC neuronal growth promoting effect of APP. The peptide can be used for
 CC increasing the memory-retention ability of a mammal, for promoting the
 CC regeneration of damaged neurons in vivo in a mammal, for treating a
 CC condition associated with cerebral deposition of amyloid beta-protein in
 CC a human patient such as Alzheimer's disease, or for treating a
 CC neurological condition. This sequence is uniquely required for the growth
 CC -promoting activity of secreted APP (695 AA) on fibroblasts. (Updated on
 CC 25-MAR-2003 to correct PN field.)
 XX

SQ Sequence 5 AA;

Query Match 100.0%; Score 15; DB 2; Length 5;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 1 RER 3

RESULT 14

AAK77510
 ID AAK77510 standard; protein; 5 AA.

AC AAK77510;

DT 27-AUG-2003 (revised)
 DT 14-APR-1996 (first entry)

DE NeuroD basic region motif in bHLH proteins.

KW NeuroD; neurogenic differentiation; neuronal growth factor;
 KW basic helix-loop-helix secondary structure; neurogenesis;
 KW non-neuronal cell differentiation; antigen; drug screening;
 KW neurodegenerative disease; traumatic injury; gene therapy.

OS Metazoa.

PN WO9530693-A1.

PD 16-NOV-1995.

PF 08-MAY-1995; 95WO-US005741.

PR 06-MAY-1994; 94US-00239228.

PA (HUTC-) HUTCHINSON CANCER RES CENT FRED.
 PA (WEIN/) WEINTRAUB N.

PI Weintraub HM, Lee JE, Hollenberg SM, Tapscott SJ;

DR WPI; 1995-404081/51.

XX Nucleic acid molecule which hybridises with a neuroD HLH domain - is used
 PT in a method for inducing differentiation of a non-neuronal cell.
 XX

PS Example 3; Page 41; 50pp; English.

XX The NRAR basic region motif of NeuroD is shared by other proteins with
 CC the basic helix-loop-helix secondary structure, and the Drosophila
 CC Daughterless (Da) and mammalian E proteins. NeuroD induces
 CC differentiation of a non-neuronal cell into a neuron. DNA encoding NeuroD
 CC may be used in the development of probes, in the construction of
 CC recombinant cell lines and transgenic animals, and in the construction of
 CC gene therapy vectors for the repair of neuronal defects resulting from
 CC traumatic injury and neurodegenerative diseases (Alzheimer's disease,
 CC Huntington's disease, Parkinson's disease). Transformed host cells are
 CC used (1) as a source of neuronal growth factors, (2) in transient and
 CC continuous cultures for anti-cancer drug screening, and (3) as sources of
 CC recombinant NeuroD for use as an antigen in diagnostic antibody
 CC production. (Updated on 27-AUG-2003 to correct OS field.)

XX Sequence 5 AA;
 SQ
 Query Match 100.0%; Score 15; DB 2; Length 5;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 3 RER 5

RESULT 15

AAW22449
 ID AAW22449 standard; peptide; 5 AA.

AC AAW22449;

DT 02-OCT-1997 (first entry)

DE NeuroD1 NRAR motif.

KW Neurogenic differentiation protein; neuroD1; transcriptional activator.

OS Mus musculus.

PN WO9716548-A1.

PD 09-MAY-1997.

PF 30-OCT-1996; 96WO-US017532.

PR 02-NOV-1995; 95US-00552142.

PA (HUTC-) HUTCHINSON CANCER RES CENT FRED.
 PA (WEIN/) WEINTRAUB N.

PI Weintraub HM, Lee JE, Hollenberg SM, Tapscott SJ;

DR WPI; 1997-272117/24.

XX Nucleic acid encoding neurogenic differentiation polypeptide - useful
 PT e.g. in regulating neuronal, endocrine and gastrointestinal development.
 XX

PS Example 3; Page 23; 81pp; English.

XX The NRAR motif (AAW22449) of mouse neurogenic differentiation protein
 CC neuroD1 (see also AAW22436) is shared by other basic-helix-loop-helix
 CC (bHLH) proteins, and the Drosophila Daughterless and mammalian E
 CC proteins. Similar motifs (see also AAW22453 and AAW22454) have been found
 CC in Drosophila Atonal and mammalian achaete-scute homologue proteins,
 CC which are thought to be involved in neurogenesis. The basic region of
 CC bHLH proteins is important for DNA binding site recognition, and there is
 CC homology between neuroD1 and other neuroproteins in this functional
 CC region
 XX

SQ Sequence 5 AA;

Query Match 100.0%; Score 15; DB 2; Length 5;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 3 RER 5

RESULT 16

AAW12517
 ID AAW12517 standard; peptide; 5 AA.

AC AAW12517;

DT 22-APR-1997 (first entry)
 XX Interleukin-6 antagonist 82.
 XX
 XX Interleukin-6; IL-6; antagonist; inhibitor; autoimmune disease; skin;
 XX intestine; systemic lupus erythematosus; chronic rheumatism.
 XX
 XX Synthetic.
 XX
 XX Key Location/Qualifiers
 XX Modified-site 5 /note= "amidated"
 XX JF08311098-A.
 XX
 XX 26-NOV-1996.
 XX
 XX 22-MAY-1995; 95JP-00146742.
 XX
 XX 22-MAY-1995; 95JP-00146742.
 XX (DAIL) DAICEL CHEM IND LTD.
 XX (FUJI) FUJISAWA PHARM CO LTD.
 XX WPI; 1997-061811/06.
 XX
 XX Interleukin-6 antagonistic peptide(a) comprising arginine - useful for
 XX treating autoimmune, renal, skin and intestinal diseases.
 XX
 XX Example 82; Page 12; 20pp; Japanese.
 XX
 XX The present peptide is a specific example of new interleukin-6
 XX antagonists of the general formula E-F-G-H-Arg-NH₂, where E, F and H each
 XX represent any optionally protected amino acid and where G is preferably
 XX an arg residue having an opt. protected guanidino group, but can be any
 XX amino acid. The peptides are useful for treating autoimmune diseases
 XX (e.g. systemic lupus erythematosus or chronic rheumatism), renal, skin
 XX and intestinal diseases
 XX
 XX Sequence 5 AA;
 XX
 XX Query Match 100.0%; Score 15; DB 2; Length 5;
 XX Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 XX Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 XX
 XX QY 1 RER 3
 XX Db |||
 XX 3 RER 5
 XX
 XX RESULT 17
 XX AAW71013
 XX ID AAW71013 standard; peptide; 5 AA.
 XX
 XX AC AAW71013;
 XX
 XX 25-MAR-2003 (revised)
 XX 21-OCT-1998 (first entry)
 XX
 XX Motif of neuroD1 and Drosophila daughterless and mammalian E proteins.
 XX
 XX Basic helix-loop-helix; bHLH; neuroD; neuroectodermal tumour;
 XX classification; medulloblastoma; Drosophila daughterless;
 XX mammalian E protein.
 XX
 XX Unidentified.
 XX
 XX US5795723-A.
 XX
 XX 18-AUG-1998.
 XX
 XX 07-AUG-1997; 97US-00910973.
 XX

PR 06-MAY-1994; 94US-00239238.
 PR 02-NOV-1995; 95US-00552142.
 PR 30-OCT-1996; 96WO-US017532.
 XX
 XX (HUTC-) HUTCHINSON CANCER RES CENT FRED.
 XX
 XX Tapscott SJ, Olson JM;
 XX WPI; 1998-466661/40.
 XX
 XX Classifying neuroectodermal tumours from expression pattern of basic-
 XX helix-loop-helix genes - especially for identifying medulla:blastoma and
 XX assessing its aggressiveness, specifically associated with expression of
 XX BHLH genes neuroD 1-3.
 XX
 XX Example 3; Col 18; 43pp; English.
 XX
 XX The present sequence represents a motif found in neuroD1 and Drosophila
 XX daughterless and mammalian E proteins. NeuroD is a member of the basic
 XX helix-loop-helix (bHLH) protein family. The bHLH genes are a family of
 XX genes associated with vertebrate neuronal, endocrinal and
 XX gastrointestinal development. The observed pattern of neuroD expression
 XX distinguishes subclasses of neuroectodermal tumours. The specification
 XX describes a method for the classification of human neuroectodermal
 XX tumours. The method comprises measuring, in a tumour sample, expression
 XX of at least one basic bHLH gene and identifying the tumour subclass by
 XX matching expression to predetermined expression profiles for known
 XX subclasses. For classifying the tumour as a medulloblastoma, the bHLH
 XX gene detected is neuroD1 and neuroD3. The method is used to classify
 XX neuroectodermal tumours, and to identify medulloblastoma and for
 XX prognosis of this as aggressive. (Updated on 25-MAR-2003 to correct PR
 XX field.)
 XX
 XX Sequence 5 AA;
 XX
 XX Query Match 100.0%; Score 15; DB 2; Length 5;
 XX Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 XX Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 XX
 XX QY 1 RER 3
 XX Db |||
 XX 3 RER 5
 XX
 XX RESULT 18
 XX AAW94160
 XX ID AAW94160 standard; peptide; 5 AA.
 XX
 XX AC AAW94160;
 XX
 XX 14-APR-1999 (first entry)
 XX
 XX BC loop sequence of fluorescein-binding monobody clone pLB24.6.
 XX
 XX Fibronectin type III; Fn3; monobody; beta-strand domain; loop region;
 XX specific binding partner; SBP; catalysis; LRS; fluorescein.
 XX
 XX Unidentified.
 XX
 XX WO9856915-A2.
 XX
 XX 17-DEC-1998.
 XX
 XX 12-JUN-1998; 98WO-US012099.
 XX
 XX 12-JUN-1997; 97US-0049410P.
 XX (RESE) RESEARCH CORP TECHNOLOGIES INC.
 XX
 XX Koide S;
 XX
 XX WPI; 1999-060331/05.
 XX

PT Fibrinectin type III (Fn3) polypeptide monobody (artificial mini-
PT antibodies) comprising Fn3 P-strand domain sequences that are linked to
PT loop region sequences, useful in therapeutic, diagnostic and catalytic
PT applications.

XX Example 12; Page 42; 96pp; English.

XX The invention relates to a synthetic fibrinectin type III (Fn3)
CC polypeptide monobody that comprises Fn3 beta-strand domain sequences that
CC are linked to loop region sequences (LRSe). One or more of the loop
CC sequences in the synthetic Fn3 vary by deletion, insertion, or
CC replacement of at least 2 amino acids from the corresponding LRSe in wild
CC type Fn3. Host cells containing an expression vector comprising the
CC synthetic Fn3 nucleic acid are used for the production of the Fn3
CC monobody. The invention also provides methods of identifying the amino
CC acid sequence of a polypeptide molecule (i) capable of binding to a
CC specific binding partner (SBP) so as to form a polypeptide:SBP complex;
CC (ii) capable of catalysing a chemical reaction with a catalysed rate
CC constant, Kcat, and an uncatalysed rate constant, Kuncat, such that the
CC ratio of the Kcat/Kuncat is greater than 10. Sequences AAW94155-63
CC represent BC loop sequences of fluorescein-binding monobody clones from
CC library #2

XX Sequence 5 AA;

Query Match 100.0%; Score 15; DB 2; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 19

AAB91809
ID AAB91809 standard; peptide; 5 AA.

XX AAB91809;
XX 22-JUN-2001 (first entry)

DE Amyloid beta-protein fragment peptide SEQ ID NO:985.

XX Protection; endogenous therapeutic peptide; peptidase; conjugation;
KW blood component; modification; succinimidyl; maleimido group; amino;
KW hydroxyl; thiol; hormone; growth factor; neurotransmitter.

OS Homo sapiens.
OS Synthetic.

PN WC200069900-A2.

XX 23-NOV-2000.

PF 17-MAY-2000; 2000WO-US013576.

XX 17-MAY-1999; 99US-0134406P.

PR 10-SEP-1999; 99US-0153406P.

PR 15-OCT-1999; 99US-0159783P.

XX (CONJ-) CONJUCHEM INC.

XX Bridon DP, Ezrin AM, Milner PG, Holmes DL, Thibaudau K;

XX WPI; 2001-112059/12.

XX Modifying and attaching therapeutic peptides to albumin prevents

PT peptidase degradation, useful for increasing length of in vivo activity.

XX Disclosure; Page 516; 733pp; English.

XX The present invention describes a modified therapeutic peptide (I)

CC comprising a therapeutically active amino acid region (III) and a
CC reactive group (II) (e.g. succinimidyl and maleimido groups) attached to
CC a less therapeutically active amino acid region (IV), which covalently
CC bonds with amino/hydroxyl/thiol groups on blood components to form a
CC peptidase stabilised therapeutic peptide composed of 3-50 amino acids.
CC (I) are useful for modifying therapeutic peptides e.g. hormones, growth
CC factors and neurotransmitters, to protect them from peptidase activity in
CC vivo for the treatment of various disorders. Endogenous therapeutic
CC peptides are not suitable as drug candidates as they require frequent
CC administration due to rapid degradation by peptidases in the body.
CC Modifying and attaching therapeutic peptides to albumin prevents or
CC reduces the action of peptidases to increase length of activity (half
CC life) and specificity as bonding to large molecules decreases
CC intracellular uptake and interference with physiological processes.
CC AAB90829 to AAB92441 represent peptides which can be used in the
CC exemplification of the present invention

XX Sequence 5 AA;

Query Match 100.0%; Score 15; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 1 RER 3

RESULT 20

AAB91776
ID AAB91776 standard; peptide; 5 AA.

XX AAB91776;

XX 22-JUN-2001 (first entry)

DE Amyloid beta-protein fragment peptide SEQ ID NO:952.

XX Protection; endogenous therapeutic peptide; peptidase; conjugation;
KW blood component; modification; succinimidyl; maleimido group; amino;
KW hydroxyl; thiol; hormone; growth factor; neurotransmitter.

OS Homo sapiens.
OS Synthetic.

PN WC200069900-A2.

XX 23-NOV-2000.

PF 17-MAY-2000; 2000WO-US013576.

XX 17-MAY-1999; 99US-0134406P.

PR 10-SEP-1999; 99US-0153406P.

PR 15-OCT-1999; 99US-0159783P.

XX (CONJ-) CONJUCHEM INC.

XX Bridon DP, Ezrin AM, Milner PG, Holmes DL, Thibaudau K;

XX WPI; 2001-112059/12.

XX Modifying and attaching therapeutic peptides to albumin prevents
PT peptidase degradation, useful for increasing length of in vivo activity.

XX Disclosure; Page 505; 733pp; English.

XX The present invention describes a modified therapeutic peptide (I)
CC comprising a therapeutically active amino acid region (III) and a
CC reactive group (II) (e.g. succinimidyl and maleimido groups) attached to
CC a less therapeutically active amino acid region (IV), which covalently
CC bonds with amino/hydroxyl/thiol groups on blood components to form a
CC peptidase stabilised therapeutic peptide composed of 3-50 amino acids.
CC (I) are useful for modifying therapeutic peptides e.g. hormones, growth

CC factors and neurotransmitters, to protect them from peptidase activity in
 CC vivo for the treatment of various disorders. Endogenous therapeutic
 CC peptides are not suitable as drug candidates as they require frequent
 CC administration due to rapid degradation by peptidases in the body.
 CC Modifying and attaching therapeutic peptides to albumin prevents or
 CC reduces the action of peptidases to increase length of activity (half
 CC life) and specificity as bonding to large molecules decreases
 CC intracellular uptake and interference with physiological processes.
 CC AAB90829 to AAB92441 represent peptides which can be used in the
 CC exemplification of the present invention

XX SQ Sequence 5 AA;

Query Match 100.0%; Score 15; DB 4; Length 5;

Best Local Similarity 100.0%; Pred. No. 1.4e+06; Indels 0; Gaps 0;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3

Db 1 RER 3

RESULT 21

AAU01260
 ID AAU01260 standard; peptide; 5 AA.

XX AC AAU01260;

XX DT 18-JUL-2001 (first entry)

XX DE B. subtilis pantothenate synthetase altered C-terminus #1.

XX KW Pantothenate synthetase; panC; pantothenate biosynthesis; NDI; NDI1;
 XX vitamin B5; nutritional supplement; panto-compound; pantoate; RBS;
 XX ribosome binding site.

XX OS Bacillus subtilis.

XX PN WC200121772-A2.

XX PD 29-MAR-2001.

XX PF 21-SEP-2000; 2000WO-US025993.

XX PR 21-SEP-1999; 99US-00400494.

XX PR 07-JUN-2000; 2000US-0210072P.

XX PR 28-JUL-2000; 2000US-0221836P.

XX PR 24-AUG-2000; 2000US-0227860P.

XX PA (ONNI-) OMNIGENE BIOPRODUCTS.

XX PI Yocum RR, Patterson TA, Hermann T, Pero JG;

XX DR WPI; 2001-218644/22.

XX DR N-PSDB; AAS01012, AAS02311.

XX PT New recombinant microorganism which overexpress a Bacillus subtilis
 PT pantothenate biosynthetic enzyme, useful for the high yield production of
 PT panto-compounds such as pantothenate and pantoate.

XX PS Disclosure; Page 49; 292pp; English.

XX CC The sequence is the C-terminus of B. subtilis pantothenate synthetase
 CC (encoded by the panC gene, an enzyme of the pantothenate biosynthetic
 CC pathway), as encoded by the artificial ribosome binding sites NDI and
 CC NDI1 used for panto, the next gene in the operon. Pantothenate, also known
 CC as vitamin B5, is used as a nutritional supplement in mammals and humans.
 CC The invention concerns methods of producing recombinant microorganisms
 CC overexpressing at least one B. subtilis pantothenate biosynthetic enzyme.
 CC The microorganisms and methods of producing them are useful for producing
 CC a panto-compound such as pantothenate or pantoate, which is a nutritional
 CC requirement for livestock and humans. The methods are also useful for the
 CC identification of pantothenate kinase modulators. Panto-compounds are

CC produced at a significantly higher yield than prior art methods and can
 CC be produced independent of the need to feed precursors which decreases
 CC expense

XX SQ Sequence 5 AA;

Query Match 100.0%; Score 15; DB 4; Length 5;

Best Local Similarity 100.0%; Pred. No. 1.4e+06; Indels 0; Gaps 0;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3

Db 2 RER 4

RESULT 22

ABB94415
 ID ABB94415 standard; peptide; 5 AA.

XX AC ABB94415;

XX DT 12-JUN-2002 (first entry)

XX DE Ubiquitin binding antibody clone pLE24-6 BC loop SEQ ID NO: 59.

XX KW Fibronectin type 3; mutant; stabilising mutation; Fn3; antibody;
 XX binding protein.

XX OS Unidentified.

XX PN WC200204523-A2.

XX PD 17-JAN-2002.

XX PF 11-JUL-2001; 2001WO-US021855.

XX PR 11-JUL-2000; 2000US-0217474P.

XX PA (RESE) RESEARCH CORP TECHNOLOGIES INC.
 XX (KOID/) KOIDE S.

XX PI Koide S;

XX DR WPI; 2002-171708/22.

XX PT New fibronectin type III molecule comprising a stabilizing mutation,
 PT useful for introducing more mutations for better functions, and in a
 PT wider range of applications.

XX PS Example 12; Page 145; 164pp; English.

XX CC The present invention relates to fibronectin type III (Fn3) molecules
 CC comprising a stabilising mutation as compared to a wild-type Fn3. Fn3 can
 CC be used as a scaffold to engineer artificial binding proteins.
 CC Modifications of the Fn3 scaffold that increase its stability are useful
 CC in that they allow the introduction of more mutations for better
 CC functions, and that these make it possible to use Fn3-based engineered
 CC proteins in a wider range of applications. The present sequence is a
 CC peptide described in the exemplification of the invention

XX SQ Sequence 5 AA;

Query Match 100.0%; Score 15; DB 5; Length 5;

Best Local Similarity 100.0%; Pred. No. 1.4e+06; Indels 0; Gaps 0;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3

Db 3 RER 5

RESULT 23

ABG71799

AG71799 standard; peptide; 5 AA.
ABG71799;
23-JAN-2003 (first entry)
bHLH family neuroD protein basic region motif.
Mouse; neuroD3; neuroD; basic-helix-loop-helix; bHLH; differentiation;
neuroD; endocrine; gastrointestinal; development; transgenic; embryo;
birth defect; spontaneous abortion; stem cell; cancer;
neural growth factor; tumor; diagnostic; motor; sensory;
traumatic neural injury; hearing; vision; brain; spinal cord;
malabsorption syndrome; gastrointestinal dysmotility syndrome;
Hirsh Prung's disease; therapeutic; DNA binding site.
Mus musculus.
US6444463-B1.
03-SEP-2002.
07-FEB-2000; 2000US-00493227.
06-MAY-1994; 94US-00239238.
08-MAY-1995; 95WO-US005741.
02-NOV-1995; 95US-00552142.
30-OCT-1996; 96WO-US017532.
07-AUG-1997; 97US-00910973.
05-AUG-1998; 98WO-US016417.
(HUTC-) HUTCHINSON CANCER RES CENT FRED.
Tapscott SJ;
WPI; 2003-056678/05.
New neurogenic differentiation gene, useful in gene therapy to correct
traumatic neural injury that has resulted in loss of motor or sensory
neural function and for constructing recombinant cell lines.
Example 3; Col 55; 43pp; English.
The invention discloses an isolated nucleic acid molecule which encodes a
functionally active human neuroD3 polypeptide. NeuroD proteins represent
a new family within the basic-helix-loop-helix (bHLH) family which are
implicated in the regulation of differentiation. NeuroD proteins are
particularly involved in neuronal, endocrine and gastrointestinal
development. The nucleic acid is useful for constructing recombinant cell
lines, transgenic embryos and animals and for quantifying the level of
expression of neuroD in a cell. Birth defects and spontaneous abortions
may result from expression of an abnormal neuroD protein. The
polynucleotide sequences permit the establishment of primary cultures of
proliferating embryonic neuronal stem cells under conditions mimicking
those that are active in development and cancer. The resultant cell lines
find use as sources of novel neural growth factors, in assays for
identifying novel neuronal growth factors which can be used for screening
anti-cancer drugs capable of driving terminal differentiation in neural
tumours, for producing antibodies useful in diagnostic assays and for
screening for compounds capable of modulating the activity of neuroD.
Transformed host cells, nucleic acids and polypeptides are also useful
for treating sites of traumatic neural injury where motor or sensory
neural activity has been lost, e.g. hearing or vision loss and brain or
spinal cord damage. The host cells find use in the treatment of
malabsorption syndromes or gastrointestinal dysmotility syndromes (Hirsh
Prung's Disease). The cell lines also find use in screening for candidate
therapeutic agents capable of either substituting for neuroD or
correcting the cellular defect caused by a defective neuroD. The sequence
presented is the bHLH family neuroD protein basic region motif which is
similar to the basic region motif of neuroD proteins and is responsible
for DNA binding site recognition
Sequence 5 AA;

Query Match 100.0%; Score 15; DB 6; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
DB 3 RER 5
RESULT 24
ABB99606
ID ABB99606 standard; peptide; 5 AA.
XX AC ABB99606;
XX DT 28-MAR-2003 (first entry)
XX DE Peptide derived from human amyloid precursor protein (APP).
XX KW Amyloid precursor protein; APP; protein derivative;
XX KW neurodegenerative disease; Alzheimer's disease; cognitive enhancer.
XX OS Synthetic.
XX OS Homo sapiens.
XX PN WO200283729-A2.
XX PD 24-OCT-2002.
XX PF 17-APR-2002; 2002WO-GB001769.
XX PR 18-APR-2001; 2001GB-00009558.
XX PR 17-AUG-2001; 2001GB-00020084.
XX PR 30-NOV-2001; 2001US-00998491.
XX PR 28-MAR-2002; 2002GB-00007387.
XX PA (UYOP-) UNIV OPEN.
XX PI Mileusnic R, Rose SPR;
XX WPI; 2003-111814/10.
Derivatives of polypeptides, useful for treating neurodegenerative
disease e.g. Alzheimer's disease, comprises one functional amino acid
residue or derivative protected by a protective group.
Claim 74; Page 65; 85pp; English.
The present sequence is derived from amyloid precursor protein (APP).
Derivatives of the invention may be based on APP sequences. The
specification describes a derivative of a polypeptide in which at least
one functional group of at least one amino acid residue or derivative is
protected by a protective group. This derivative is of the formula given
in ABB99625. The derivative is useful in medicine and in the preparation
of a medicament for use in the treatment of a neurodegenerative disease
e.g. Alzheimer's disease. It is also useful as a cognitive enhancer
Sequence 5 AA;
Query Match 100.0%; Score 15; DB 6; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
DB 1 RER 3
RESULT 25
ABB99607
ID ABB99607 standard; peptide; 5 AA.
XX

AC ABB99607;
 XX 28-MAR-2003 (first entry)
 XX Peptide derived from human amyloid precursor protein (APP).
 XX
 XX Amyloid precursor protein; APP; protein derivative;
 KW neurodegenerative disease; Alzheimer's disease; cognitive enhancer.
 XX
 XX Synthetic.
 OS Homo sapiens.
 XX WO2002083729-A2.
 XX
 XX 24-OCT-2002.
 XX
 XX 17-APR-2002; 2002WO-GB001769.
 XX
 XX 18-APR-2001; 2001GB-00009558.
 PR 17-AUG-2001; 2001GB-00020084.
 PR 30-NOV-2001; 2001US-00998491.
 PR 28-MAR-2002; 2002GB-00007387.
 XX
 XX (UYOP-) UNIV OPEN.
 XX
 XX Mileusnic R, Rose SPR;
 PI WPI; 2003-111814/10.
 DR
 XX
 XX Derivatives of polypeptides, useful for treating neurodegenerative
 PT disease e.g. Alzheimer's disease, comprises one functional amino acid
 PT residue or derivative protected by a protective group.
 XX
 XX Claim 74; Page 65; 85pp; English.
 PS
 XX The present sequence is derived from amyloid precursor protein (APP).
 CC Derivatives of the invention may be based on APP sequences. The
 CC specification describes a derivative of a polypeptide in which at least
 CC one functional group of at least one amino acid residue or derivative is
 CC protected by a protective group. This derivative is of the formula given
 CC in ABB99625. The derivative is useful in medicine and in the preparation
 CC of a medicament for use in the treatment of a neurodegenerative disease
 CC e.g. Alzheimer's disease. It is also useful as a cognitive enhancer
 XX
 XX Sequence 5 AA;
 SQ
 Query Match 100.0%; Score 15; DB 6; Length 5;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 RER 3
 Db |||
 3 RER 5
 RESULT 26
 ASU62548
 ID ABU62548 standard; peptide; 5 AA.
 XX
 AC ABU62548;
 XX
 XX 18-SEP-2003 (first entry)
 DT
 DE Human secreted amyloid precursor protein alpha (sAPP alpha) region.
 XX
 XX Human; secreted amyloid precursor protein; sAPP; sAPP alpha;
 KW inflammation; ApoE3; apolipoprotein E; Alzheimer's disease; epilepsy;
 KW traumatic brain injury; stroke; antinflammatory; anticonvulsant;
 KW cerebroprotective; vulnerary; tranquiliser; nootropic; neuroprotective.
 XX
 XX Homo sapiens.
 OS
 XX US2003069198-A1.
 PN

XX 10-APR-2003.
 PD
 XX 10-JUN-2002; 2002US-00166482.
 PF
 XX 28-AUG-1998; 98US-00141951.
 PR
 XX (BARG/) BARGER S W.
 PA
 XX Barger SW;
 PI WPI; 2003-540888/51.
 DR
 XX Reducing inflammation caused by secreted amyloid precursor protein (sAPP)
 PT in brain of a mammal, by administering a compound which inhibits amino
 PT terminal region of sAPP involved in inflammatory response.
 XX
 XX Example 5; Page 5; 9pp; English.
 PS
 XX The invention relates to a method for reducing inflammation caused by
 CC secreted amyloid precursor protein (sAPP) in the brain of a mammal or
 CC potentiating the neuroprotective effect of sAPP alpha in a person,
 CC involving administering a compound which inhibits the amino terminal
 CC region of sAPP or sAPP alpha, respectively involved in the inflammatory
 CC response. The method is useful for reducing inflammation caused by sAPP
 CC in the brain of a mammal, preferably human, or for potentiating the
 CC neuroprotective effect of sAPP alpha in a person. The method is
 CC preferably useful for reducing inflammation due to reduced levels of
 CC ApoE3 or inflammation caused by Alzheimer's disease or traumatic brain
 CC injury. The method is useful for treating epilepsy, stroke, traumatic
 CC brain injury and Alzheimer's disease. This sequence represents a peptide
 CC region of human sAPP alpha
 XX
 XX Sequence 5 AA;
 SQ
 Query Match 100.0%; Score 15; DB 6; Length 5;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 RER 3
 Db |||
 1 RER 3
 RESULT 27
 AAR62104
 ID AAR62104 standard; peptide; 6 AA.
 XX
 AC AAR62104;
 XX
 XX 25-MAR-2003 (revised)
 DT 27-APR-1995 (first entry)
 DE
 XX Hydrophilic motif from nuclear protein antigens.
 XX
 KW Small ribonucleoprotein complex; U1 snRNP; 70K protein; epitope;
 KW autoantibody; immunoinfective cluster virus; nuclear protein antigen;
 KW systemic rheumatic disorder; human immunodeficiency virus; HIV-1;
 KW centromere CENP-B; thyroglobulin-h; thyroid peroxidase; scleroderma;
 KW systemic lupus erythematosus.
 XX
 XX Homo sapiens.
 OS
 XX WO9420141-A1.
 PN
 XX 15-SEP-1994.
 PD
 XX 10-MAR-1994; 94WC-US002631.
 PF
 XX 11-MAR-1993; 93US-00029850.
 PR
 XX (UYSC-) UNIV SOUTHERN CALIFORNIA.
 PA
 XX

PI Douvas A, Takehana Y, Ehresmann G;
 XX MPI; 1994-302689/37.
 XX
 PT Methods for treating immunoinfective cluster virus infections - utilise
 PT antibodies or fragments characteristic of auto antibodies produced by
 PT patients with rheumatic disorders.
 XX
 PS Disclosure; Page 8; 106pp; English.
 XX
 CC This sequence is an example of an alternating acidic/basic amino acid,
 CC hydrophilic motif possibly found in nuclear protein antigens. As well as
 CC occurring in normal human proteins, the motif is found in similar form in
 CC immunoinfective cluster viruses. The motif serves as an epitope for anti-
 CC viral antibodies and also for autoantibodies which occur in high titre in
 CC patients suffering from systemic rheumatic disorders. Sera from such
 CC patients could be used for treatment of immunoinfective cluster virus
 CC (e.g. HIV, EBV, rubella virus) infections. (Updated on 25-MAR-2003 to
 CC correct PN field.)
 XX
 SQ Sequence 6 AA;
 Query Match 100.0%; Score 15; DB 2; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 RER 3
 DB 2 RER 4
 RESULT 28
 AAR62189
 ID AAR62189 standard; protein; 6 AA.
 XX
 AC AAR62189;
 XX
 XX 25-MAR-2003 (revised)
 DT 03-MAY-1995 (first entry)
 XX
 DE U1 snRNP 70K protein amino acids 471-476, homologous to EBV motif.
 XX
 KW Small ribonucleoprotein complex; U1 snRNP; 70K protein; epitope;
 KW antibody; immunoinfective cluster virus; nuclear protein antigen;
 KW systemic rheumatic disorder; Epstein-Barr virus; EBV na protein;
 KW systemic lupus erythematosus; scleroderma.
 XX
 OS Homo sapiens.
 XX
 PN WO9420141-A1.
 XX
 PD 15-SEP-1994.
 XX
 PF 10-MAR-1994; 94WO-US002631.
 XX
 PR 11-MAR-1993; 93US-00029850.
 XX
 PA (UYSC-) UNIV SOUTHERN CALIFORNIA.
 XX
 PI Douvas A, Takehana Y, Ehresmann G;
 XX MPI; 1994-302689/37.
 XX
 PT Methods for treating immunoinfective cluster virus infections - utilise
 PT antibodies or fragments characteristic of auto antibodies produced by
 PT patients with rheumatic disorders.
 XX
 PS Disclosure; Page 69; 106pp; English.
 XX
 CC A comparison of the U1 snRNP 70K protein sequence with proteins from
 CC immunoinfective cluster viruses revealed widespread homologies. The
 CC importance of these homologous motifs is that they are epitopes for
 CC autoantibodies occurring in high titres in systemic rheumatic disorders.

CC Sera from such patients could be used for treatment of immunoinfective
 CC cluster virus infections. (Updated on 25-MAR-2003 to correct PN field.)
 XX
 SQ Sequence 6 AA;
 Query Match 100.0%; Score 15; DB 2; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 RER 3
 DB 2 RER 4
 RESULT 29
 AAW21203
 ID AAW21203 standard; peptide; 6 AA.
 XX
 AC AAW21203;
 XX
 XX 29-JUL-1997 (first entry)
 DT
 XX
 DE Farnesyl synthetase derived signal oligopeptide #3.
 XX
 KW Hydrophilic; signal oligopeptide; hydrophilicity maxima; vaccine; HIV;
 KW competitive inhibitor; feedback regulator; synthesis; gastrin precursor;
 KW charge; polarity; farnesyl synthetase; plasminogen activator inhibitor 1;
 KW hydroxymethylglutaryl coenzyme A reductase; glucagon precursor; rhesus;
 KW gonadoliberin precursor; plasminogen activator inhibitor 2; prolamin;
 KW Alzheimer amyloid A4; corticotropin releasing factor binding protein;
 KW apolipoprotein E; herpes virus 1 glycoprotein B; HSV1; human; OMVVS;
 KW herpes virus 2 glycoprotein B; HSV2; collagenase; apolipoprotein A;
 KW Treponema pallidum membrane protein; TWPA; islet amyloid polypeptide;
 KW fibroblast MMP1; schistosoma elastase precursor; schistosomin;
 KW hepatitis delta antigen; rev protein; HIV; VILV; angiotensinogen.
 XX
 OS Homo sapiens.
 XX
 PN WO9519568-A1.
 XX
 PD 20-JUL-1995.
 XX
 PF 12-JAN-1995; 95WO-US000575.
 XX
 PR 14-JAN-1994; 94US-00182248.
 XX
 PA (RATH/) RATH M.
 XX
 PI Rath M;
 XX
 DR WPI; 1995-263953/34.
 XX
 PT Identifying signal oligopeptide(s) in protein sequence(s) - shown as
 PT regions of max. hydrophilicity, used in modulating communication between
 PT protein(s).
 XX
 PS Claim 5; Page 23; 88pp; English.
 XX
 CC The sequences given in AAW21201-560 represent hydrophilic signal oligo-
 CC peptides. These signal oligopeptides are localised on the surface of the
 CC protein and are represented by the hydrophilicity maxima of the protein.
 CC These peptides are enriched in charged amino acids arranged with neutral
 CC spacer amino acids. The specific signal character of these oligopeptides
 CC is determined by a characteristic combination of conformation and charge
 CC within the signal sequence. These oligopeptides may be used as vaccines
 CC in the treatment of human disease, as competitive inhibitors to prevent
 CC or reduce the metabolic action or interaction of a selected protein by
 CC blocking its specific signal sequences, or as therapeutic agents to
 CC function as feedback regulators to reduce synthesis rate of a selected
 CC protein. These peptides may be modified by omitting one or more amino
 CC acids at the N- and/or C-terminal, by substituting one or more amino
 CC acids without consideration of charge and polarity, by substituting one
 CC or more amino acids with amino acid residues with similar charge and/or

CC polarity, by omitting one or more amino acids or a combination of these
 XX
 SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 2; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 4 RER 6

RESULT 30
 AAW21037
 ID AAW21037 standard; peptide; 6 AA.
 XX
 AC AAW21037;
 XX
 DT 19-JUN-1997 (first entry)
 XX
 DE Lipolytic enzyme opt. N- or C-terminal extension peptide #5.
 XX
 KW Lipolytic enzyme; detergent; lard; cotton swatch; laundry; dishwashing.
 XX
 OS Synthetic.
 XX
 DN WC9707202-A1.
 XX
 PD 27-FEB-1997.
 XX
 PF 12-AUG-1996; 96WO-DK000341.
 XX
 PR 11-AUG-1995; 95DK-00000905.
 PR 29-SEP-1995; 95DK-00001096.
 PR 14-FEB-1996; 96US-0011627P.
 PR 01-APR-1996; 96DK-00000374.
 PR 07-MAY-1996; 96US-0016754P.
 XX
 PA (NOVO) NOVO-NORDISK AS.

PI Okkels JS, Svendsen A, Borch K, Thellersen M, Patkar SA;
 PI Petersen DA, Royer JC, Kretzschmar T;
 PI WPI; 1997-165287/15.
 DR
 XX
 PT Lipolytic enzyme with high capacity to remove lard in one wash cycle -
 PT also related DNA, vectors and transformed cells, useful in laundry and
 PT dishwashing formulations.
 PS Claim 17; Page 244; 274pp; English.
 XX

CC The sequences given in AAW21033-92 are peptides which may be added to the
 CC N- or C-terminal of the lipolytic enzyme of the invention. The lipolytic
 CC enzyme, when present in a specified detergent composition, is able to
 CC remove at least 15% more lard from soiled cotton swatches (9 by 9 cm)
 CC than an equiv. enzyme-free compn. in a one-cycle wash assay. The assay
 CC uses 7 lard-stained cotton swatches in 1000 ml water (3.2 mM Ca2+/Mg2+;
 CC ratio 5:1; 5 g/l detergent; pH 10 plus 12500 IU of enzyme/l) for 20 min
 CC at 30 deg.C, in a thermostated Terg-O-Meter, then 15 min rinsing,
 CC drying overnight and Soxhlet extrn. and quantification of fatty material.
 CC The enzyme may be used in laundry and dishwashing formulations. It is are
 CC able to remove a substantial amount of lard in a single cycle under
 CC realistic washing conditions

SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 2; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||

Db 4 RER 6
 RESULT 31
 AAW23160
 ID AAW23160 standard; peptide; 6 AA.
 XX
 AC AAW23160;
 XX
 DT 28-OCT-1997 (first entry)
 XX
 DE Terminal peptide extension for lipolytic enzyme.

XX
 KW Lipolytic enzyme; modification; peptide extension; detergent;
 KW washing powder; dishwashing composition; pitch removal; paper; pulp;
 KW manufacture; degreasing; hide; sheepskin; wool; catalysis;
 KW organic synthesis; transesterification; esterification; ester hydrolysis;
 KW baking; defatting.

XX
 OS Synthetic.

XX
 PN WC9704078-A1.

XX
 PD 06-FEB-1997.

XX
 PF 12-JUL-1996; 96WO-DK000321.

XX
 PR 14-JUL-1995; 95DK-00000832.

XX
 PR 13-SEP-1995; 95DK-00001013.

XX
 PR 29-SEP-1995; 95DK-00001096.

XX
 PR 21-NOV-1995; 95DK-00001306.

XX
 PR 14-FEB-1996; 96US-0011634P.

XX
 PR 01-APR-1996; 96DK-00000372.

XX
 PR 07-MAY-1996; 96US-0020461P.

XX
 PA (NOVO) NOVO-NORDISK AS.

XX
 PI Fuglsang CC, Okkels JS, Pertersen DA, Patkar SA, Thellersen M;
 PI Vind J, Jorgensen ST;

XX
 WPI; 1997-132621/12.

XX
 PT Modified lipolytic enzymes with peptide extensions at one or both ends -
 PT esp. for use in detergent and dishwashing compsn., have improved
 PT substrate affinity, stability and wash performance.

XX
 PS Claim 12; Page 178; 197pp; English.

XX
 CC A lipolytic enzyme, modified by a peptide extension, e.g. the present
 CC sequence, of its carboxy and/or amino terminus, can be used in
 CC detergents, particularly in washing powders or dishwashing compositions.
 CC It may also be used to remove pitch in paper and pulp manufacture, to
 CC degrease hides, sheepskins and wool, to catalyse organic synthesis, e.g.
 CC (trans)esterification or ester hydrolysis, in baking and in other
 CC degreasing/defatting processes. The peptide extension(s) increases
 CC substrate affinity, confers stability and especially improves wash
 CC performance, i.e. better lipid soil removal, reducing the amount of
 CC enzyme used

XX
 SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 2; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 4 RER 6

RESULT 32
 AAY55251
 ID AAY55251 standard; peptide; 6 AA.

XX AC AAY55251;
 XX DT 07-JAN-2000 (first entry)
 XX DE ATCC HB 11885 monoclonal antibody 9079 releasing peptide SEQ ID NO:145.
 XX KW Antibody releasing peptide; CD34; hybridoma; binding; antigen;
 KW cell surface antigen; identification; haematopoietic stem cell; tumour;
 KW cancer; immune system; therapy; displacement.
 XX OS Synthetic.
 OS Homo sapiens.
 XX PN US5968753-A.
 XX PD 19-OCT-1999.
 XX PF 07-JUN-1995; 95US-00482228.
 XX PR 14-JUN-1994; 94US-00259427.
 XX PA (NEXE-) NEXELL THERAPEUTICS INC.
 XX PI Guillermo R, Helgerson SL, Deans RJ, Tseng-Law J, Kobori JA;
 PI Al-Abdaly FA;
 XX WPI; 1999-590399/50.
 XX PT Short peptides useful for displacing antibodies from cell surface
 PT antigens.
 XX PS Example 9; Col 32; 81pp; English.
 XX CC The present invention describes peptides of 4-17 amino acids which
 CC displace either the anti-CD34 monoclonal antibody designated 561, the
 CC anti-CD34 mouse monoclonal antibody produced by the hybridoma ATCC HB-
 CC 11646 (designated 9069), the anti-CD34 antibody produced by hybridoma
 CC ATCC HB-11885 (9079), or the anti-human breast cancer antibody produced
 CC by hybridoma ATCC HB-11884 (9187), from a cell surface antigen on a
 CC target cell. The peptides are useful for displacing antibodies bound to
 CC cell surfaces to release cells that have been positively selected by
 CC antibody-mediated binding to beads or other solid support. AAY55107 to
 CC AAY55319 represent peptides used in the exemplification of the present
 CC invention
 XX SQ Sequence 6 AA;
 Query Match 100.0%; Score 15; DB 2; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 RER 3
 DB 1 RER 3
 RESULT 33
 AAY86997
 ID AAY86997 standard; peptide; 6 AA.
 XX AC AAY86997;
 XX DT 09-MAY-2000 (first entry)
 XX DE Human haematopoietic CD34+ cell binding peptide SEQ ID #145.
 XX KW Human; haematopoietic CD34+ cell; binding peptide; monoclonal antibody;
 KW non-enzymatic cell selection method; haematopoietic stem cell;
 KW haematopoietic progenitor cell; antibody 561; breast cancer cell;
 KW antibody 9187; cell surface determinant; diagnostic cell based assay.
 XX OS Homo sapiens.

XX PN US6017719-A.
 XX PD 25-JAN-2000.
 XX PF 07-JUN-1995; 95US-00482528.
 XX PR 14-JUN-1994; 94US-00259427.
 XX PA (NEXE-) NEXELL THERAPEUTICS INC.
 XX PI Guillermo R, Helgerson SL, Deans RJ, Tseng-Law J, Kobori JA;
 PI Al-Abdaly FA;
 XX WPI; 2000-136676/12.
 XX PT Non-enzymatic method for the positive selection of target cells from a
 PT heterogeneous cell suspension, useful for selecting human breast cancer
 PT cells from a patient's blood or bone marrow.
 XX PS Example 9; Col 36; 82pp; English.
 XX CC This sequence represents a human haematopoietic CD34+ cell binding
 CC peptide, and was used to test the method of the invention. The method is
 CC a non-enzymatic method for the positive selection of one or more target
 CC cells from a heterogeneous cell suspension, by using specific peptides
 CC which effect the displacement and release of a specific target cell from
 CC a specific monoclonal antibody. The method is useful for positive
 CC selection and specific release of target human haematopoietic
 CC stem/progenitor cells bound by the monoclonal anti-CD34 antibodies and
 CC the antibody 561. The method is also useful for positive selection and
 CC specific release of target human breast cancer cells, bound by the
 CC monoclonal anti-breast cancer antibody 9187, from a patient's blood or
 CC bone marrow. Identification of peptide epitopes for antibodies which
 CC recognise cell surface determinants also allows construction of
 CC diagnostic cell based assays. The peptide mediated release is enzyme free
 CC and thus leaves the cell surface proteins intact. Moreover, peptide
 CC mediated release leaves the target cell free of bound antibody or
 CC antibody fragments. The method also produces a high yield of functional
 CC target cells and is relatively inexpensive to carry out
 XX SQ Sequence 6 AA;
 Query Match 100.0%; Score 15; DB 3; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 RER 3
 DB 1 RER 3
 RESULT 34
 AAB36760
 ID AAB36760 standard; peptide; 6 AA.
 XX AC AAB36760;
 XX DT 16-FEB-2001 (first entry)
 XX DE HRG-beta1 library B variant 12.
 XX KW Heregulin; ErbB receptor; transplantation; cancer;
 KW nervous system disease; musculature; epithelium.
 XX OS Unidentified.
 XX PN US6136558-A.
 XX PD 24-OCT-2000.
 XX PF 09-FEB-1998; 98US-00020880.

PR 10-FEB-1997; 97US-0037581P.
XX (GETH) GENENTECH INC.
XX Jones JT, Fairbrother WJ, Ballinger MD, Wells JA, Sliwkowski MK;
XX WPI; 2000-678767/66.
XX
XX New variants of heregulin, useful e.g. for treating cancer, comprises
PT specific amino acid alterations that increase affinity for ErbB
PT receptors.
XX
XX Example 3; Col 73; 58pp; English.
XX
XX The present invention relates to variants of heregulin that can bind to
CC an ErbB receptor and include a portion of the 175-230 region of native
CC human heregulin-beta1. The variants may be used to promote ex vivo
CC survival, proliferation and differentiation of cells, particularly when
CC intended for transplantation. They may also be used to treat a wide range
CC of cancers and diseases of the nervous system, musculature and epithelium
XX
XX Sequence 6 AA:
SQ
Query Match 100.0%; Score 15; DB 3; Length 6;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 RER 3
Db 2 RER 4
RESULT 35
AA94683
ID AA94683 standard; protein; 6 AA.
XX
XX
XX AA94683;
XX
XX 01-DEC-2000 (first entry)
XX Human zsig83 hydrophilic region peptide.
XX
XX Alpha-helical protein; zsig83; cell growth; differentiation; cancer;
XX proliferation; chromosome 22q13.1-q13.2; cytostatic; vulneray;
XX degenerative condition; metastasis; wound healing.
XX
XX Homo sapiens.
XX
XX WO2000050594-A2.
XX
XX 31-AUG-2000.
XX
XX 25-FEB-2000; 2000WO-US004816.
XX
XX 26-FEB-1999; 99US-00259131.
XX
XX (ZYMO) ZYMOGENETICS INC.
XX
XX Presnell SR;
XX
XX WPI; 2000-572091/53.
XX
XX Alpha-helical protein zsig83, its antibodies and the polynucleotide
PT encoding the protein useful for treating disorders associated with
PT abnormal cell growth e.g. cancer and agonists useful for treating wounds.
XX
XX Disclosure; Page 77; 83pp; English.
XX
XX This invention relates to a novel human alpha-helical protein designated
CC zsig83. Zsig83 plays a role in the process of cell growth,
CC differentiation, or proliferation. The zsig83 gene is located on
CC chromosome 22 at position 22q13.1-q13.2. Included in the invention are
CC polynucleotide sequences encoding the zsig83 protein, expression vectors

CC containing the zsig83 DNA sequence, a cultured cell containing the
CC expression vector, and antibodies specific to the zsig83 protein. The
CC zsig83 protein contains 5 alpha helix regions (represented by sequences
CC AA94677-Y94681) and also contains epitope bearing regions (represented
CC by sequences AA94688-Y94698) to which the antibodies are directed. The
CC protein exhibits cytostatic and vulneray activity. The zsig83 protein and
CC nucleotide sequences and antibodies are used for treating disorders
CC associated with abnormal cell growth e.g. cancer, degenerative conditions
CC and metastasis. The zsig83 protein and its agonists or antagonists are
CC useful for promoting wound healing. The zsig83 DNA sequence can be used
CC to identify defective zsig83 genes and may therefore be used as a
CC diagnostic indicator of cancer. The present sequence represents a
CC hydrophilic region of the human zsig83 protein
XX
XX Sequence 6 AA;
SQ
Query Match 100.0%; Score 15; DB 3; Length 6;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 RER 3
Db 4 RER 6
RESULT 36
AA97624
ID AA97624 standard; peptide; 6 AA.
XX
XX
XX AA97624;
XX
XX 21-SEP-2001 (first entry)
XX
XX Neuropeptide Y (NPY) modulator, Ac-Ile-Trp-Arg-Glu-Arg-Tyr-NH2.
XX
XX Neuropeptide Y; NPY modulator; agonist; antagonist; appetite; obesity;
XX eating disorder; blood pressure; cardiovascular response; hypertension;
XX libido; sexual dysfunction; circadian rhythm; sleep disorder;
XX gastrointestinal disorder; gallbladder disorder;
XX central nervous system disorder; insulin-related disorder;
XX type II diabetes; pain; anorectic; hypertensive; hypotensive; cardiant;
XX vasotropic; analgesic.
XX
XX Synthetic.
XX
XX Key Location/Qualifiers
FH Modified-site 1 /note= "N-terminal acetyl"
FT Modified-site 6 /note= "C-terminal amide"
FT
FT
XX US6235718-B1.
XX
XX 22-MAY-2001.
XX
XX 02-DEC-1999; 99US-00449914.
XX
XX 09-AUG-1996; 96US-0023588P.
XX 07-AUG-1997; 97US-00907408.
XX
XX (UYCI-) UNIV CININNATI.
XX
XX Balasubramaniam A, Chance WT;
XX
XX WPI; 2001-440207/47.
XX
XX New tripeptide derivatives are neuropeptide Y modulators useful for
PT controlling appetite, blood pressure, cardiovascular response, libido and
PT circadian rhythm.
XX
XX Disclosure; Col 19; 13pp; English.
XX
XX The invention relates to neuropeptide Y (NPY) agonist and antagonist
CC

CC peptides. The peptides comprise a unit of 3 amino acids (N-terminally
 CC designated A1, A2 and A3, where A1 is joined to two chemical groups,
 CC designated R1 and R2, and A3 is linked to a chemical moiety designated W.
 CC These are defined as follows: A1 is Trp (or a derivative thereof, e.g.,
 CC Tcc), Gln (or a derivative thereof), a tethered amino acid with an indole
 CC ring (e.g., NMe-Trp), Phe, Hyp, Pyr, Bch, Nal, Tcc, Asn, Nva, Abu, Tyr,
 CC Tic-OH, Phe, Tip or Dip; A2 is Arg, N-Me-Arg, C-alpha-Me-Arg, Orn, Cit,
 CC HArg(R)2 (where R is selected from the group consisting of H, alkyl,
 CC aralkyl, or alkylaryl), or Lys-epsilon-NH2; A3 is N-Me-Tyr, C-alpha-Me-
 CC Tyr, Tic-OH, Tic-Dip, Trp, Phe, des-carboxylic-Tyr or Tyr-R' (where R'
 CC is H or a lipophilic group); R1 and R2 are independently H, 1-12C alkyl, 6
 CC -18C aryl, 1-18C acyl, 7-18C aralkyl, 7-18C alkaryl or
 CC dihydrotrigonellinate; and W is OH, NR3R4 or OR5 (where R3, R4 and R5 are
 CC independently H, 1-12C alkyl, 6-18C aryl, 1-12C acyl, 7-18C aralkyl or 7-
 CC 18C alkaryl). The peptide bonds may be replaced by pseudo-peptide bonds.
 CC In certain embodiments of the invention, A1 or A3 are absent, in which
 CC case the N-terminal residue is joined to groups R1 and R2, and the C-
 CC terminal residue is joined to group W. The invention also encompasses
 CC peptide dimers, in which peptides of the invention are dimerised with
 CC cysteine, dicarboxylic acids, or diaminodicarboxylic acids. The invention
 CC further encompasses peptides of the formula Ac-[A1-A2-A3]n-NH2, where n
 CC is 1, 2, or 3, or a cyclic peptide of formula cyclo-[A1-A2-A3], cyclo-[A1-
 CC A2-A3]2, or cyclo-[A1-A2-A3-A2-A1]. The neuropeptide Y agonists and
 CC antagonists of the invention are useful for regulating appetite, blood
 CC pressure, cardiovascular response, libido and circadian rhythm, and may
 CC therefore be used in the treatment of obesity, eating disorders,
 CC hypertension, alterations in sexual function, and sleep disorders. They
 CC may also be used in the treatment of gastrointestinal disorders
 CC (including gallbladder disorders), central nervous system disorders,
 CC insulin-related disorders (e.g., type II diabetes), and pain. The present
 CC sequence represents a neuropeptide Y modulator peptide of the invention
 CC
 XX Sequence 6 AA;

Query Match 100.0%; Score 15; DB 4; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 ||||
 Db 3 RER 5

RESULT 37
 AAB82171
 ID AAB82171 standard; peptide; 6 AA.

XX AAB82171;
 AC AAB82171;
 XX
 XX 20-JUL-2001 (first entry)
 DT
 XX Peptide #22 used in a method for inducing Th1 immune response.
 DE
 XX Antibacterial; antiallergic; cytostatic; Th1 immune response inducer;
 KW vaccine; infectious disease; allergy; cancer.
 XX
 XX Synthetic.
 OS
 XX W0200126682-A2.
 FN
 XX 19-APR-2001.
 PD
 XX 13-OCT-2000; 2000WO-US028443.
 PF
 XX 14-OCT-1999; 99GB-00024351.
 PR
 XX (DOWC) DOW CHEM CO.
 PA Brennan F;
 PT Increasing the level of Th1-type responses to molecules, used to treat

PT infectious diseases, allergies and cancer, comprises conjugating the
 PT molecule to a plant virus.

XX Example 11; Page 70; 89pp; English.

XX The present invention relates to a method for increasing the level of a
 CC Th1-type immune response to a molecule. The method comprising conjugating
 CC the molecule to a heterologous peptide expressed by a plant virus, and
 CC administering the conjugate to an animal. The present sequence is one
 CC such heterologous peptide, which may be used in the method of the present
 CC invention. The method is useful for treating infectious diseases,
 CC allergies and cancer, by administering appropriate antigens conjugated to
 CC a plant virus

XX Sequence 6 AA;

Query Match 100.0%; Score 15; DB 4; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 ||||
 Db 1 RER 3

RESULT 38
 ABB99622
 ID ABB99622 standard; peptide; 6 AA.

XX ABB99622;
 AC ABB99622;
 XX
 XX 28-MAR-2003 (first entry)
 DT
 XX Peptide derived from human amyloid precursor protein (APP).
 DE
 XX Amyloid precursor protein; APP; protein derivative;
 KW neurodegenerative disease; Alzheimer's disease; cognitive enhancer.
 XX
 XX Synthetic.
 OS
 XX Homo sapiens.
 OS
 XX W0200283729-A2.
 FN
 XX 24-OCT-2002.
 PD
 XX 17-APR-2002; 2002WO-GB001769.
 PF
 XX 18-APR-2001; 2001GB-00009558.
 PR
 XX 17-AUG-2001; 2001GB-00020084.
 PR
 XX 30-NOV-2001; 2001US-0098491.
 PR
 XX 28-MAR-2002; 2002GB-00007387.
 PR
 XX (UYOP-) UNIV OPEN.
 PA
 XX Mileusenic R, Rose SPR;
 PI
 XX WPI, 2003-111814/10.
 DR
 XX
 XX Disclosures; Page 9; 85pp; English.

PT Derivatives of polypeptides, useful for treating neurodegenerative
 PT disease e.g. Alzheimer's disease, comprises one functional amino acid
 PT residue or derivative protected by a protective group.
 XX
 XX Disclosures; Page 9; 85pp; English.
 CC The present sequence is derived from amyloid precursor protein (APP).
 CC Derivatives of the invention are based on APP sequences. The
 CC specification describes a derivative of a polypeptide in which at least
 CC one functional group of at least one amino acid residue or derivative is
 CC protected by a protective group. This derivative is of the formula given
 CC in ABB99625. The derivative is useful in medicine and in the preparation
 CC of a medicament for use in the treatment of a neurodegenerative disease
 CC e.g. Alzheimer's disease. It is also useful as a cognitive enhancer

SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 6; Length 6;
Best Local Similarity 100.0%; Pred. NO. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
|||
Db 2 RER 4

RESULT 39

ADE65222
ID ADE65222 standard; peptide; 6 AA.

AC ADE65222;

DT 29-JAN-2004 (first entry)

DE Corticotropin-releasing factor-2 polypeptide, SEQ ID NO 525.

XX corticotropin-releasing factor-2; CRF2; myopathic; osteopathic;
XX hypotensive; cardiant; vasotropic; antimigraine; cerebroprotective;
XX neotropic; neuroprotective; anorectic; antidiabetic; analgesic;
XX antiallergic; tranquilizer; anxiolytic; antidepressant; antiarthritic;
XX gene therapy.

XX Unidentified.

OS WC2003062277-A1.

XX 31-JUL-2003.

XX 16-JAN-2003; 2003WO-US001454.

XX 16-JAN-2002; 2002US-0349117P.

XX 29-APR-2002; 2002US-0376337P.

XX 14-JUN-2002; 2002US-0388895P.

XX 19-SEP-2002; 2002US-0411988P.

XX (PROC) PROCTER & GAMBLE CO.

XX Isfort RJ, Mazur WA;

XX WPI; 2003-787975/74.

XX New non-native peptide derived from corticotropin-releasing factor-2,
XX useful for treatment and prevention of e.g. muscular atrophy, also
XX related nucleic acid and antibodies.

XX Example 2; SEQ ID NO 525; 304pp; English.

XX The invention relates to a novel non-native peptide derived from
XX corticotropin-releasing factor-2 (CRF2). The CRF2 peptides have the
XX following activities: myopathic, osteopathic, hypotensive, cardiant,
XX vasotropic, antimigraine, cerebroprotective, neotropic, neuroprotective,
XX anorectic, antidiabetic, analgesic, antiallergic, tranquilizer,
XX anxiolytic, antidepressant, and antiarthritic. The CRF2 peptides, and
XX related compounds derived from other proteins, are used to prevent or
XX treat disorders modulated by the CRF2 receptor, e.g. skeletal muscle
XX atrophy or wasting, and bone disorders, however caused; heart/circulatory
XX diseases (e.g. hypertension, congestive heart failure, heart attack,
XX reperfusion injury, migraine, stroke, memory loss, Alzheimer's disease,
XX dementia); joint disorders (osteoarthritis or rheumatoid arthritis);
XX metabolic disease (obesity or diabetes); pain; allergy; stress; anxiety;
XX low levels of adrenocorticotrophic hormone; anorexia nervosa; depression;
XX function. Nucleic acids, optionally labelled, that encode the CRF2
XX peptides are used as primers and probes for amplification, also for gene
XX synthesis and for recombinant production of CRF2 peptides, including use
XX in gene therapy. Antibodies specific for the CRF2 peptides are used to
XX evaluate expression of the CRF2 peptides after gene therapy. This
XX sequence represents a novel native CRF polypeptide of the invention.

XX SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 7; Length 6;
Best Local Similarity 100.0%; Pred. NO. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
|||
Db 3 RER 5

RESULT 40

ADE65221
ID ADE65221 standard; peptide; 6 AA.

XX ADE65221;

XX 29-JAN-2004 (first entry)

XX Corticotropin-releasing factor-2 polypeptide, SEQ ID NO 524.

XX corticotropin-releasing factor-2; CRF2; myopathic; osteopathic;
XX hypotensive; cardiant; vasotropic; antimigraine; cerebroprotective;
XX neotropic; neuroprotective; anorectic; antidiabetic; analgesic;
XX antiallergic; tranquilizer; anxiolytic; antidepressant; antiarthritic;
XX gene therapy.

XX Unidentified.

OS WO2003062277-A1.

XX 31-JUL-2003.

XX 16-JAN-2003; 2003WO-US001454.

XX 16-JAN-2002; 2002US-0349117P.

XX 29-APR-2002; 2002US-0376337P.

XX 14-JUN-2002; 2002US-0388895P.

XX 19-SEP-2002; 2002US-0411988P.

XX (PROC) PROCTER & GAMBLE CO.

XX Isfort RJ, Mazur WA;

XX WPI; 2003-787975/74.

XX New non-native peptide derived from corticotropin-releasing factor-2,
XX useful for treatment and prevention of e.g. muscular atrophy, also
XX related nucleic acid and antibodies.

XX Example 2; SEQ ID NO 524; 304pp; English.

XX The invention relates to a novel non-native peptide derived from
XX corticotropin-releasing factor-2 (CRF2). The CRF2 peptides have the
XX following activities: myopathic, osteopathic, hypotensive, cardiant,
XX vasotropic, antimigraine, cerebroprotective, neotropic, neuroprotective,
XX anorectic, antidiabetic, analgesic, antiallergic, tranquilizer,
XX anxiolytic, antidepressant, and antiarthritic. The CRF2 peptides, and
XX related compounds derived from other proteins, are used to prevent or
XX treat disorders modulated by the CRF2 receptor, e.g. skeletal muscle
XX atrophy or wasting, and bone disorders, however caused; heart/circulatory
XX diseases (e.g. hypertension, congestive heart failure, heart attack,
XX reperfusion injury, migraine, stroke, memory loss, Alzheimer's disease,
XX dementia); joint disorders (osteoarthritis or rheumatoid arthritis);
XX metabolic disease (obesity or diabetes); pain; allergy; stress; anxiety;
XX low levels of adrenocorticotrophic hormone; anorexia nervosa; depression;
XX also to reduce body temperature and to control appetite or cognitive
XX function. Nucleic acids, optionally labelled, that encode the CRF2
XX peptides are used as primers and probes for amplification, also for gene
XX synthesis and for recombinant production of CRF2 peptides, including use
XX in gene therapy. Antibodies specific for the CRF2 peptides are used to
XX evaluate expression of the CRF2 peptides after gene therapy. This

CC sequence represents a novel native CRF polypeptide of the invention.
 XX
 SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 7; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 3 RER 5

RESULT 41

AD65159
 ID ADE65159 standard; peptide; 6 AA.

XX AC ADE65159;

XX DT 29-JAN-2004 (first entry)

XX DE Corticotropin-releasing factor-2 polypeptide, SEQ ID NO 462.

XX KW corticotropin-releasing factor-2; CRF2; myopathic; osteopathic;
 KW hypotensive; cardiant; vasotropic; antimigraine; cerebroprotective;
 KW neurotropic; neuroprotective; anorectic; antidiabetic; analgesic;
 KW antiallergic; tranquilizer; anxiolytic; antidepressant; antiarthritic;
 KW gene therapy.

XX OS Unidentified.

XX PN W02003062277-A1.

XX PD 31-JUL-2003.

XX PF 16-JAN-2003; 2003WO-US001454.

XX PR 16-JAN-2002; 2002US-0349117P.

XX PR 29-APR-2002; 2002US-0376337P.

XX PR 14-JUN-2002; 2002US-038895P.

XX PR 19-SEP-2002; 2002US-0411988P.

XX PA (PROC) PROCTER & GAMBLE CO.

XX PI Isfort RJ, Mazur WA;

XX DR WPI; 2003-787975/74.

XX PT New non-native peptide derived from corticotropin-releasing factor-2,

XX PT useful for treatment and prevention of e.g. muscular atrophy, also

XX PT related nucleic acid and antibodies.

XX PS Example 2; SEQ ID NO 462; 304pp; English.

XX CC The invention relates to a novel non-native peptide derived from
 CC corticotropin-releasing factor-2 (CRF2). The CRF2 peptides have the
 CC following activities: myopathic, osteopathic, hypotensive, cardiant,
 CC vasotropic, antimigraine, cerebroprotective, neurotropic, neuroprotective,
 CC anorectic, antidiabetic, analgesic, antiallergic, tranquilizer,
 CC anxiolytic, antidepressant, and antiarthritic. The CRF2 peptides, and
 CC related compounds derived from other proteins, are used to prevent or
 CC treat disorders modulated by the CRF2 receptor, e.g. skeletal muscle
 CC atrophy or wasting, and bone disorders, however caused; heart/circulatory
 CC diseases (e.g. hypertension, congestive heart failure, heart attack,
 CC reperfusion injury, migraine, stroke, memory loss, Alzheimer's disease,
 CC dementia); joint disorders (osteoarthritis or rheumatoid arthritis);
 CC metabolic disease (obesity or diabetes); pain; allergy; stress; anxiety;
 CC low levels of adrenocorticotrophic hormone; anorexia nervosa; depression;
 CC also to reduce body temperature and to control appetite or cognitive
 CC function. Nucleic acids, optionally labelled, that encode the CRF2
 CC peptides are used as primers and probes for amplification, also for gene
 CC synthesis and for recombinant production of CRF2 peptides, including use
 CC in gene therapy. Antibodies specific for the CRF2 peptides are used to

CC evaluate expression of the CRF2 peptides after gene therapy. This
 CC sequence represents a novel native CRF polypeptide of the invention.
 XX
 SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 7; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 3 RER 5

RESULT 42

ADE51507
 ID ADE51507 standard; peptide; 6 AA.

XX AC ADE51507;

XX DT 29-JAN-2004 (first entry)

XX DE CRF2 non-native polypeptide, SEQ ID NO 524.

XX KW non-native; corticotropin-releasing factor-2; CRF2; myopathic;
 KW osteopathic; hypotensive; cardiant; vasotropic; antimigraine;
 KW cerebroprotective; neurotropic; neuroprotective; anorectic; antidiabetic;
 KW analgesic; antiallergic; tranquilizer; anxiolytic; antidepressant;
 KW antiarthritic.

XX OS Unidentified.

XX PN W02003062268-A2.

XX PD 31-JUL-2003.

XX PF 16-JAN-2003; 2003WO-US001451.

XX PR 16-JAN-2002; 2002US-0349117P.

XX PR 29-APR-2002; 2002US-0376337P.

XX PR 14-JUN-2002; 2002US-038895P.

XX PR 19-SEP-2002; 2002US-0411988P.

XX PA (PROC) PROCTER & GAMBLE CO.

XX PI Isfort RJ, Mazur WA;

XX DR WPI; 2003-787974/74.

XX PT New non-native peptide derived from corticotropin-releasing factor-2,

XX PT useful for treatment and prevention of e.g. muscular atrophy, also

XX PT related nucleic acid and antibodies.

XX PS Example 2; SEQ ID NO 524; 300pp; English.

XX CC The invention relates to a novel non-native peptide derived from
 CC corticotropin-releasing factor-2 (CRF2). The non-native CRF2 peptides
 CC have the following activities: myopathic, osteopathic, hypotensive,
 CC cardiant, vasotropic, antimigraine, cerebroprotective, neurotropic,
 CC neuroprotective, anorectic, antidiabetic, analgesic, antiallergic,
 CC tranquilizer, anxiolytic, antidepressant, and antiarthritic. The non-
 CC native CRF2 peptides, and related compounds derived from other proteins,
 CC are used to prevent or treat disorders modulated by the CRF2 receptor,
 CC e.g. skeletal muscle atrophy or wasting, and bone disorders, however
 CC caused; heart/circulatory diseases (e.g. hypertension, congestive heart
 CC failure, heart attack, reperfusion injury, migraine, stroke, memory loss,
 CC Alzheimer's diseases, dementia); joint disorders (osteoarthritis or
 CC rheumatoid arthritis); metabolic disease (obesity or diabetes); pain;
 CC allergy; stress; anxiety; low levels of adrenocorticotrophic hormone;
 CC anorexia nervosa; depression; also to reduce body temperature and to
 CC control appetite or cognitive function. Nucleic acids, optionally
 CC labelled, that encode the non-native CRF2 peptides are used as primers
 CC and probes for amplification, also for gene synthesis and for recombinant

CC production of the non-native CRF2 peptides, including use in gene
 CC therapy. Antibodies specific for the non-native CRF2 peptides are used to
 CC evaluate expression of the non-native CRF2 peptides after gene therapy.
 CC This sequence represents a CRF2 non-native polypeptide of the invention.

XX SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 7; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 3 RER 5

RESULT 43

AD5E1445
 ID ADE51445 standard; peptide; 6 AA.

XX AC ADE51445;

XX DT 29-JAN-2004 (first entry)

XX CRF2 non-native polypeptide, SEQ ID NO 462.

XX non-native; corticotropin-releasing factor-2; CRF2; myopathic;
 KW osteopathic; hypotensive; cardiant; vasotropic; antimigraine;
 KW cerebroprotective; nootropic; neuroprotective; anorectic; antidiabetic;
 KW analgesic; antiallergic; tranquilizer; anxiolytic; antidepressant;
 KW antiarthritic.

XX Unidentified.

XX WO2003062268-A2.

XX PN 31-JUL-2003.

XX 16-JAN-2003; 2003WO-US001451.

XX 16-JAN-2002; 2002US-0349117P.

XX 29-APR-2002; 2002US-0376337P.

XX 14-JUN-2002; 2002US-0388895P.

XX 19-SEP-2002; 2002US-0411988P.

XX (PROC) PROCTER & GAMBLE CO.

XX Isfort RJ, Mazur WA;

XX WPI; 2003-787974/74.

XX New non-native peptide derived from corticotropin-releasing factor-2,
 PT useful for treatment and prevention of e.g. muscular atrophy, also
 PT related nucleic acid and antibodies.

XX Example 2; SEQ ID NO 462; 300pp; English.

XX The invention relates to a novel non-native peptide derived from
 CC corticotropin-releasing factor-2 (CRF2). The non-native CRF2 peptides
 CC have the following activities: myopathic, osteopathic, hypotensive,
 CC cardiant, vasotropic, antimigraine, cerebroprotective, nootropic,
 CC neuroprotective, anorectic, antidiabetic, analgesic, antiallergic,
 CC tranquilizer, anxiolytic, antidepressant, and antiarthritic. The non-
 CC native CRF2 peptides, and related compounds derived from other proteins,
 CC are used to prevent or treat disorders modulated by the CRF2 receptor,
 CC e.g. skeletal muscle atrophy or wasting, and bone disorders, however
 CC caused; heart/circulatory diseases (e.g. hypertension, congestive heart
 CC failure, heart attack, reperfusion injury, migraine, stroke, memory loss,
 CC Alzheimer's diseases, dementia); joint disorders (osteoarthritis or
 CC rheumatoid arthritis); metabolic disease (obesity or diabetes); pain;
 CC allergy; stress; anxiety; low levels of adrenocorticotrophic hormone;
 CC anorexia nervosa; depression; also to reduce body temperature and to
 CC control appetite or cognitive function. Nucleic acids, optionally

CC labelled, that encode the non-native CRF2 peptides are used as primers
 CC and probes for amplification, also for gene synthesis and for recombinant
 CC production of the non-native CRF2 peptides, including use in gene
 CC therapy. Antibodies specific for the non-native CRF2 peptides are used to
 CC evaluate expression of the non-native CRF2 peptides after gene therapy.
 CC This sequence represents a CRF2 non-native polypeptide of the invention.

XX SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 7; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 |||
 Db 3 RER 5

RESULT 44

AD5E1508

XX ID ADE51508 standard; peptide; 6 AA.

XX AC ADE51508;

XX DT 29-JAN-2004 (first entry)

XX CRF2 non-native polypeptide, SEQ ID NO 525.

XX non-native; corticotropin-releasing factor-2; CRF2; myopathic;
 KW osteopathic; hypotensive; cardiant; vasotropic; antimigraine;
 KW cerebroprotective; nootropic; neuroprotective; anorectic; antidiabetic;
 KW analgesic; antiallergic; tranquilizer; anxiolytic; antidepressant;
 KW antiarthritic.

XX Unidentified.

XX WO2003062268-A2.

XX PN 31-JUL-2003.

XX 16-JAN-2003; 2003WO-US001451.

XX 16-JAN-2002; 2002US-0349117P.

XX 29-APR-2002; 2002US-0376337P.

XX 14-JUN-2002; 2002US-0388895P.

XX 19-SEP-2002; 2002US-0411988P.

XX (PROC) PROCTER & GAMBLE CO.

XX Isfort RJ, Mazur WA;

XX WPI; 2003-787974/74.

XX New non-native peptide derived from corticotropin-releasing factor-2,
 PT useful for treatment and prevention of e.g. muscular atrophy, also
 PT related nucleic acid and antibodies.

XX Example 2; SEQ ID NO 525; 300pp; English.

XX The invention relates to a novel non-native peptide derived from
 CC corticotropin-releasing factor-2 (CRF2). The non-native CRF2 peptides
 CC have the following activities: myopathic, osteopathic, hypotensive,
 CC cardiant, vasotropic, antimigraine, cerebroprotective, nootropic,
 CC neuroprotective, anorectic, antidiabetic, analgesic, antiallergic,
 CC tranquilizer, anxiolytic, antidepressant, and antiarthritic. The non-
 CC native CRF2 peptides, and related compounds derived from other proteins,
 CC are used to prevent or treat disorders modulated by the CRF2 receptor,
 CC e.g. skeletal muscle atrophy or wasting, and bone disorders, however
 CC caused; heart/circulatory diseases (e.g. hypertension, congestive heart
 CC failure, heart attack, reperfusion injury, migraine, stroke, memory loss,
 CC Alzheimer's diseases, dementia); joint disorders (osteoarthritis or
 CC rheumatoid arthritis); metabolic disease (obesity or diabetes); pain;
 CC allergy; stress; anxiety; low levels of adrenocorticotrophic hormone;
 CC control appetite or cognitive function. Nucleic acids, optionally

CC anorexia nervosa; depression; also to reduce body temperature and to
 CC control appetite or cognitive function. Nucleic acids, optionally
 CC labelled, that encode the non-native CRF2 peptides are used as primers
 CC and probes for amplification, also for gene synthesis and for recombinant
 CC production of the non-native CRF2 peptides, including use in gene
 CC therapy. Antibodies specific for the non-native CRF2 peptides are used to
 CC evaluate expression of the non-native CRF2 peptides after gene therapy.
 CC This sequence represents a CRF2 non-native polypeptide of the invention.
 XX
 XX SQ Sequence 6 AA;

Query Match 100.0%; Score 15; DB 7; Length 6;
 Best Local Similarity 100.0%; Pred. No. 1.4e+06;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
 |||
 Db 3 RER 5

RESULT 45

AA24590
 ID AAR24590 standard; peptide; 7 AA.

AC AAR24590;

XX 25-MAR-2003 (revised)

DT 03-DEC-1992 (first entry)

XX Immunomodulatory peptide.

XX Immunodeficiencies; immunosuppression; T-cell subset; immunotherapy;

XX inflammation; wounds; lymphocyte; vaccine.

XX Synthetic.

XX WC9209628-A1.

XX 11-JUN-1992.

XX 22-NOV-1991; 91WO-US008795.

XX 23-NOV-1990; 90US-00617494.

XX (IMMU-) IMMUNODYNAMICS INC.

XX Atkin A;

XX WPI; 1992-217021/26.

XX New synthetic immunomodulatory peptide(s) - for treating

PT immunodeficiencies, immunosuppression and T-cell subset deviations and

PT immuno-therapy of infections, inflammation, wounds etc.

XX Claim 9; Page 34; 52pp; English.

XX The immunomodulatory peptide is a specific example of a peptide cpd. (or

CC an acid or base salt) constructed by combination and/or overlapping of

CC the amino acid sequences AIB1XB2A2, ABE3XA4B4, BSA5XA6B6, B7A7XB8A8,

CC A9B9, A10A11, B10A12, and B11B12 (X= Ala, Gly, Ile, Leu, Phe or Val, A1-

CC A12 each= Arg, Asn, Gln, Lys, Phe or Val; B1-B12 each= Asp, Glu, Tyr, Phe

CC or Val. The synthetic peptide may be used for immunomodulation of various

CC immunodeficiencies and immunosuppressed conditions, T-cell subset and

CC lymphocyte deviations, enhancement of a vaccine's efficacy, as well as for

CC immunotherapy, including infections, local or systemic complications of

CC non-infectious diseases, postoperative inflammations, wounds and burns.

CC See also AAR24583-R24701. (Updated on 25-MAR-2003 to correct FN field.)

XX SQ Sequence 7 AA;

Query Match 100.0%; Score 15; DB 2; Length 7;

Best Local Similarity 100.0%; Pred. No. 1.4e+06;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
 |||
 Db 1 RER 3

RESULT 46

AAW21423
 ID AAW21423 standard; peptide; 7 AA.

XX AC AAW21423;

XX 29-JUL-1997 (first entry)

XX Alzheimer amyloid A4 derived signal oligopeptide #9.

XX Hydrophilic; signal oligopeptide; hydrophilicity maxima; vaccine; SIV;
 KW competitive inhibitor; feedback regulator; synthesis; gastrin precursor;
 KW charge; polarity; farnesyl synthetase; plasminogen activator inhibitor 1;
 KW hydroxymethylglutaryl coenzyme A reductase; glucagon precursor; rhesus;
 KW gonadoliberin precursor; plasminogen activator inhibitor 2; prorenin;
 KW Alzheimer amyloid A4; corticotropin releasing factor binding protein;
 KW apolipoprotein E; herpes virus 1 glycoprotein B; HSV1; human; OMV5;
 KW herpes virus 2 glycoprotein B; HSV2; collagenase; apolipoprotein A;
 KW Treponema pallidum membrane protein; TWPA; islet amyloid polypeptide;
 KW fibroblast MWPI; schistosoma elastase precursor; schistosomin;
 KW hepatitis delta antigen; rev protein; HIV; VILV; angiotensinogen.

OS Homo sapiens.

XX WC9519568-A1.

XX 20-JUL-1995.

XX 12-JAN-1995; 95WO-US000575.

XX 14-JAN-1994; 94US-00182248.

XX (RATH/) RATH M.

XX Rath M;

XX WPI; 1995-263953/34.

XX Identifying signal oligopeptide(s) in protein sequence(s) - shown as

PT regions of max. hydrophilicity, used in modulating communication between

PT protein(s).

XX Claim 5; Page 60; 88pp; English.

XX The sequences given in AAW21201-560 represent hydrophilic signal oligo-

CC peptides. These signal oligopeptides are localised on the surface of the

CC protein and are represented by the hydrophilicity maxima of the protein.

CC These peptides are enriched in charged amino acids arranged with neutral

CC spacer amino acids. The specific signal character of these oligopeptides

CC is determined by a characteristic combination of conformation and charge

CC within the signal sequence. These oligopeptides may be used as vaccines

CC in the treatment of human disease, as competitive inhibitors to prevent

CC or reduce the metabolic action or interaction of a selected protein by

CC blocking its specific signal sequences, or as therapeutic agents to

CC function as feedback regulators to reduce synthesis rate of a selected

CC protein. These peptides may be modified by omitting one or more amino

CC acids at the N- and/or C-terminal, by substituting one or more amino

CC acids without consideration of charge and polarity, by substituting one

CC or more amino acids with amino acid residues with similar charge and/or

CC polarity, by omitting one or more amino acids or a combination of these

XX Sequence 7 AA;

Query Match 100.0%; Score 15; DB 2; Length 7;

Best Local Similarity 100.0%; Pred. No. 1.4e+06;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 5 RER 7

RESULT 47
AAR77509
ID AAR77509 standard; protein; 7 AA.
XX AC AAR77509;
XX 27-AUG-2003 (revised)
DT 14-APR-1996 (first entry)
XX DE Basic region motif in MASH neurogenesis protein.
XX NeuroD; neurogenic differentiation; neuronal growth factor;
KW basic helix-loop-helix secondary structure; neurogenesis;
KW non-neuronal cell differentiation; antigen; drug screening;
KW neurodegenerative disease; traumatic injury; gene therapy.
XX OS Mammalia.
XX WO9530693-A1.
PN 16-NOV-1995.
XX 08-MAY-1995; 95WO-US005741.
PF 06-MAY-1994; 94US-00239228.
PR (HUTC-) HUTCHINSON CANCER RES CENT FRED.
PA (WEIN/) WEINTRAUB N.
XX Weintraub HM, Lee JE, Hollenberg SM, Tapscott SJ;
XX WPI; 1995-404081/51.
XX Nucleic acid molecule which hybridises with a neuroD HLH domain - is used
PT in a method for inducing differentiation of a non-neuronal cell.
XX Example 3; Page 41; 50pp; English.
XX The basic region motif of the MASH (mammalian achaete-scute homologue),
CC neurogenesis protein is similar to that of the Drosophila Atonal protein
CC (see AAR77508) and to the basic region motif of murine NeuroD (AAR77504).
CC NeuroD induces differentiation of a non-neuronal cell into a neuron. DNA
CC encoding NeuroD may be used in the development of probes, in the
CC construction of recombinant cell lines and transgenic animals, and in the
CC resulting from traumatic injury and neurodegenerative diseases
CC (Alzheimer's disease, Huntington's disease, Parkinson's disease).
CC Transformed host cells are used (1) as a source of neuronal growth
CC factors, (2) in transient and continuous cultures for anti-cancer drug
CC screening, and (3) as sources of recombinant NeuroD for use as an antigen
CC in diagnostic antibody production. (Updated on 27-AUG-2003 to correct OS
XX field.)

XX Sequence 7 AA;
Query Match 100.0%; Score 15; DB 2; Length 7;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 48
AAR77508
ID AAR77508 standard; protein; 7 AA.
XX

AC AAR77508;
XX 14-APR-1996 (first entry)
DT XX
DE Basic region motif in Atonal neurogenesis protein.
XX NeuroD; neurogenic differentiation; neuronal growth factor;
KW basic helix-loop-helix secondary structure; neurogenesis;
KW non-neuronal cell differentiation; antigen; drug screening;
KW neurodegenerative disease; traumatic injury; gene therapy.
XX Drosophila sp.
XX WO9530693-A1.
PN 16-NOV-1995.
PD 08-MAY-1995; 95WO-US005741.
PF 06-MAY-1994; 94US-00239228.
PR (HUTC-) HUTCHINSON CANCER RES CENT FRED.
PA (WEIN/) WEINTRAUB N.
XX Weintraub HM, Lee JE, Hollenberg SM, Tapscott SJ;
XX WPI; 1995-404081/51.
XX Nucleic acid molecule which hybridises with a neuroD HLH domain - is used
PT in a method for inducing differentiation of a non-neuronal cell.
XX Example 3; Page 41; 50pp; English.
XX The basic region motif of the Drosophila Atonal protein is similar to
CC that of the MASH (mammalian achaete-scute homologue) (see AAR77509) and
CC to the basic region motif of the murine NeuroD (see AAR77504). NeuroD
CC induces differentiation of a non-neuronal cell into a neuron. DNA
CC encoding NeuroD may be used in the development of probes, in the
CC construction of recombinant cell lines and transgenic animals, and in the
CC resulting from traumatic injury and neurodegenerative diseases
CC (Alzheimer's disease, Huntington's disease, Parkinson's disease).
CC Transformed host cells are used (1) as a source of neuronal growth
CC factors, (2) in transient and continuous cultures for anti-cancer drug
CC screening, and (3) as sources of recombinant NeuroD for use as an antigen
CC in diagnostic antibody production
XX Sequence 7 AA;
Query Match 100.0%; Score 15; DB 2; Length 7;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 49
AAR77508
ID AAR77508 standard; peptide; 7 AA.
XX AC AAR77508;
XX 27-AUG-2003 (revised)
DT 02-OCT-1997 (first entry)
XX DE NARERR motif of mammalian achaete-scute homologue protein.
XX Neurogenic differentiation protein; neuroD1; neurogenesis;
KW transcriptional activator; mammalian achaete-scute homologue; MASH.
XX Unidentified.

```

XX PN W09716548-A1.
XX PD 09-MAY-1997.
XX PF 30-OCT-1996; 96WO-US017532.
XX PR 02-NOV-1995; 95US-00552142.
XX PA (HUTC-) HUTCHINSON CANCER RES CENT FRED.
XX PA (WEIN/) WEINTRAUB N.
XX PI Weintraub HM, Lee JE, Hollenberg SM, Tapscott SJ;
XX DR WPI; 1997-272117/24.
XX PT Nucleic acid encoding neurogenic differentiation polypeptide - useful
XX PT e.g. in regulating neuronal, endocrine and gastrointestinal development.
XX FS Example 3; Page 23; 81pp; English.
XX SS The NARERRR (AAW22453) and NERERRR (AAW22454) motifs are found in the
CC Drosophila Atonal and mammalian achaete-scute homologue proteins,
CC respectively, and are thought to be involved in neurogenesis. The related
CC NARER motif (AAW22449) of mouse neurogenic differentiation protein
CC neuroD1 (see also AAW22436) is shared by other basic-helix-loop-helix
CC (bHLH) proteins, and the Drosophila Daughtersless and mammalian E
CC proteins. The basic region of bHLH proteins is important for DNA binding
CC site recognition, and there is homology between neuroD1 and other
CC neuroproteins in this functional region. (Updated on 27-AUG-2003 to
CC correct OS field.)
XX SQ Sequence 7 AA;

Query Match 100.0%; Score 15; DB 2; Length 7;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

RESULT 50
AAW22453
ID AAW22453 standard; peptide; 7 AA.
XX AC AAW22453;
XX DT 02-OCT-1997 (first entry)
XX DE NARERR motif of Drosophila Atonal.
XX KW Neurogenic differentiation protein; neuroD1; neurogenesis;
XX KW transcriptional activator; Atonal.
XX OS Drosophila sp.
XX PN W09716548-A1.
XX PD 09-MAY-1997.
XX PF 30-OCT-1996; 96WO-US017532.
XX PR 02-NOV-1995; 95US-00552142.
XX PA (HUTC-) HUTCHINSON CANCER RES CENT FRED.
XX PA (WEIN/) WEINTRAUB N.
XX PI Weintraub HM, Lee JE, Hollenberg SM, Tapscott SJ;
XX DR WPI; 1997-272117/24.
XX PT Nucleic acid encoding neurogenic differentiation polypeptide - useful
XX PT e.g. in regulating neuronal, endocrine and gastrointestinal development.
XX FS Example 3; Page 23; 81pp; English.
XX SS The NARERRR (AAW22453) and NERERRR (AAW22454) motifs are found in the
CC Drosophila Atonal and mammalian achaete-scute homologue proteins,
CC respectively, and are thought to be involved in neurogenesis. The related
CC NARER motif (AAW22449) of mouse neurogenic differentiation protein
CC neuroD1 (see also AAW22436) is shared by other basic-helix-loop-helix
CC (bHLH) proteins, and the Drosophila Daughtersless and mammalian E
CC proteins. The basic region of bHLH proteins is important for DNA binding
CC site recognition, and there is homology between neuroD1 and other
CC neuroproteins in this functional region. (Updated on 27-AUG-2003 to
CC correct OS field.)
XX SQ Sequence 7 AA;

Query Match 100.0%; Score 15; DB 2; Length 7;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 3 RER 5

Search completed: March 5, 2004, 16:09:35
Job time : 72 secs

```

```

PT Nucleic acid encoding neurogenic differentiation polypeptide - useful
PT e.g. in regulating neuronal, endocrine and gastrointestinal development.
XX FS Example 3; Page 23; 81pp; English.
XX SS The NARERRR (AAW22453) and NERERRR (AAW22454) motifs are found in the
CC Drosophila Atonal and mammalian achaete-scute homologue proteins,
CC respectively, and are thought to be involved in neurogenesis. The related
CC NARER motif (AAW22449) of mouse neurogenic differentiation protein
CC neuroD1 (see also AAW22436) is shared by other basic-helix-loop-helix
CC (bHLH) proteins, and the Drosophila Daughtersless and mammalian E
CC proteins. The basic region of bHLH proteins is important for DNA binding
CC site recognition, and there is homology between neuroD1 and other
CC neuroproteins in this functional region.
XX SQ Sequence 7 AA;

```

```

Query Match 100.0%; Score 15; DB 2; Length 7;
Best Local Similarity 100.0%; Pred. No. 1.4e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 RER 3
DB 3 RER 5

```

```

Search completed: March 5, 2004, 16:09:35
Job time : 72 secs

```

GenCore version 5.1.6
Copyright (c) 1993 - 2004 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: March 5, 2004, 16:08:22 ; Search time 23 Seconds
(without alignments)
6.734 Million cell used

Title: US-09-998-491-9
Perfect score: 15
Sequence: 1 RER 3

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 389414 seqs, 51625971 residues

Total number of hits satisfying chosen parameters: 234

```
Minimum DB seq length: 0
Maximum DB seq length: 20
```

Post-processing:	Minimum Match	100%
	Maximum Match	100%
	Listing first	1000

```
Database : Issued Patents AA:
1: /cgn2_6/ptodata/
2: /cgn2_6/ptodata/
3: /cgn2_6/ptodata/
4: /cgn2_6/ptodata/
5: /cgn2_6/ptodata/
6: /cgn2_6/ptodata/
```

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query #	Match	Length	DB	ID	Description
1	15	100.0	4	3	US-08-864-301-1		Sequence 1, Appl
2	15	100.0	4	3	US-09-142-078-29		Sequence 29, Appl
3	15	100.0	4	3	US-09-357-141-29		Sequence 29, Appl
4	15	100.0	4	4	US-09-533-889-29		Sequence 29, Appl
5	15	100.0	4	4	US-09-142-080-29		Sequence 29, Appl
6	15	100.0	4	5	PCF-US92-09070-14		Sequence 14, Appl
7	15	100.0	5	1	US-08-552-142A-7		Sequence 7, Appl
8	15	100.0	5	1	US-08-704-170-13		Sequence 13, Appl
9	15	100.0	5	1	US-08-704-170-14		Sequence 14, Appl
10	15	100.0	5	1	US-08-704-170-60		Sequence 60, Appl
11	15	100.0	5	4	US-08-910-973-7		Sequence 7, Appl
12	15	100.0	5	4	US-09-499-227-7		Sequence 7, Appl
13	15	100.0	5	4	US-09-638-202A-59		Sequence 59, Appl
14	15	100.0	5	4	US-09-036-749A-59		Sequence 59, Appl
15	15	100.0	5	5	PCF-US92-09070-1		Sequence 1, Appl
16	15	100.0	5	5	PCF-US94-02631-13		Sequence 13, Appl
17	15	100.0	5	5	PCF-US94-02631-14		Sequence 14, Appl
18	15	100.0	5	5	PCF-US94-02631-60		Sequence 60, Appl
19	15	100.0	5	5	PCF-US95-05741-7		Sequence 7, Appl
20	15	100.0	6	1	US-08-704-170-4		Sequence 4, Appl
21	15	100.0	6	1	US-08-704-170-97		Sequence 97, Appl
22	15	100.0	6	2	US-08-482-228-145		Sequence 145, App
23	15	100.0	6	3	US-08-482-528-145		Sequence 145, App
24	15	100.0	6	3	US-09-020-880-60		Sequence 60, Appl
25	15	100.0	6	4	US-09-101-544-60		Sequence 60, Appl
26	15	100.0	6	4	US-09-007-288B-21		Sequence 21, Appl
27	15	100.0	6	4	US-08-932-411A-29		Sequence 29, Appl

101	15	100.0	11	4	US-09-410-551B-57	Sequence 57, Appl	174	15	100.0	17	1	US-08-071-811A-3	Sequence 3, Appl
102	15	100.0	11	4	US-08-445-638-122	Sequence 122, App	175	15	100.0	17	1	US-08-071-811A-4	Sequence 4, Appl
103	15	100.0	11	5	PCT-US91-08328-14	Sequence 14, Appl	176	15	100.0	17	1	US-08-071-811A-5	Sequence 5, Appl
104	15	100.0	11	5	PCT-US91-08328-16	Sequence 16, Appl	177	15	100.0	17	1	US-08-460-874A-22	Sequence 22, Appl
105	15	100.0	11	5	PCT-US91-08328-18	Sequence 18, Appl	178	15	100.0	17	1	US-08-460-874A-27	Sequence 27, Appl
106	15	100.0	11	5	PCT-US91-08328-20	Sequence 20, Appl	179	15	100.0	17	2	US-08-388-883B-22	Sequence 22, Appl
107	15	100.0	11	5	PCT-US92-09070-10	Sequence 10, Appl	180	15	100.0	17	2	US-08-388-883B-27	Sequence 27, Appl
108	15	100.0	12	1	US-08-704-170-2	Sequence 2, Appl	181	15	100.0	17	2	US-08-598-873-50	Sequence 50, Appl
109	15	100.0	12	3	US-09-058-459-32	Sequence 32, Appl	182	15	100.0	17	3	US-07-808-452-22	Sequence 22, Appl
110	15	100.0	12	3	US-09-058-459-33	Sequence 33, Appl	183	15	100.0	17	3	US-08-462-211A-22	Sequence 22, Appl
111	15	100.0	12	3	US-09-127-926-32	Sequence 32, Appl	184	15	100.0	17	3	US-08-462-211A-27	Sequence 27, Appl
112	15	100.0	12	3	US-09-127-926-33	Sequence 33, Appl	185	15	100.0	17	3	US-08-605-430-50	Sequence 50, Appl
113	15	100.0	12	3	US-08-083-9450-10	Sequence 10, Appl	186	15	100.0	17	4	US-09-227-357-349	Sequence 349, App
114	15	100.0	12	3	US-08-079-814-1	Sequence 1, Appl	187	15	100.0	17	4	US-09-518-036-22	Sequence 22, Appl
115	15	100.0	12	4	US-08-561-578C-24	Sequence 24, Appl	188	15	100.0	17	4	US-09-079-030-189	Sequence 189, App
116	15	100.0	12	4	US-09-347-926-11	Sequence 11, Appl	189	15	100.0	17	4	US-10-279-130-22	Sequence 22, Appl
117	15	100.0	12	4	US-09-638-202A-118	Sequence 118, App	190	15	100.0	17	4	US-09-564-677A-18	Sequence 18, Appl
118	15	100.0	12	4	US-09-526-193A-63	Sequence 63, Appl	191	15	100.0	17	5	PCT-US92-09070-8	Sequence 8, Appl
119	15	100.0	12	4	US-09-526-193A-64	Sequence 64, Appl	192	15	100.0	17	5	PCT-US92-09070-17	Sequence 17, Appl
120	15	100.0	12	4	US-09-096-749A-118	Sequence 118, App	193	15	100.0	17	5	PCT-US92-10770-22	Sequence 22, Appl
121	15	100.0	12	5	PCT-US94-06631-2	Sequence 2, Appl	194	15	100.0	18	1	PCT-US94-06192-1	Sequence 1, Appl
122	15	100.0	12	5	PCT-US94-07107A-10	Sequence 10, Appl	195	15	100.0	18	1	US-08-240-514-47	Sequence 47, Appl
123	15	100.0	13	3	US-08-388-353-634	Sequence 634, App	196	15	100.0	18	2	US-08-612-302A-47	Sequence 47, Appl
124	15	100.0	13	3	US-08-488-551B-634	Sequence 634, App	197	15	100.0	18	2	US-08-824-151-5	Sequence 5, Appl
125	15	100.0	13	3	US-08-147-592A-41	Sequence 41, Appl	198	15	100.0	18	2	US-08-902-623-8	Sequence 8, Appl
126	15	100.0	13	4	US-08-292-694A-41	Sequence 46, Appl	199	15	100.0	18	2	US-09-017-205-23	Sequence 23, Appl
127	15	100.0	13	4	US-08-292-694A-46	Sequence 4, Appl	200	15	100.0	18	3	US-08-943-173-4	Sequence 4, Appl
128	15	100.0	13	4	US-08-258-851-4	Sequence 4, Appl	201	15	100.0	18	3	US-08-943-173-5	Sequence 5, Appl
129	15	100.0	14	2	US-08-484-905-55	Sequence 55, Appl	202	15	100.0	18	4	US-09-325-601-8	Sequence 8, Appl
130	15	100.0	14	3	US-08-370-476-55	Sequence 55, App	203	15	100.0	18	4	US-09-440-781-72	Sequence 72, Appl
131	15	100.0	14	3	US-08-074-575-55	Sequence 55, App	204	15	100.0	18	4	US-09-079-030-182	Sequence 182, App
132	15	100.0	14	4	US-08-248-588-99	Sequence 99, Appl	205	15	100.0	18	4	US-09-564-677A-17	Sequence 17, Appl
133	15	100.0	14	4	US-08-992-877-35	Sequence 35, Appl	206	15	100.0	18	5	PCT-US94-01234-9	Sequence 9, Appl
134	15	100.0	14	4	US-09-638-202A-1	Sequence 1, Appl	207	15	100.0	18	5	PCT-US95-07543-4	Sequence 4, Appl
135	15	100.0	14	4	US-09-096-749A-1	Sequence 1, Appl	208	15	100.0	19	1	US-08-019-073-30	Sequence 30, Appl
136	15	100.0	14	5	PCT-US95-13975-10	Sequence 10, Appl	209	15	100.0	19	2	US-08-442-461D-15	Sequence 15, Appl
137	15	100.0	14	5	US-08-660A-1	Sequence 1, Appl	210	15	100.0	19	4	US-09-082-358B-69	Sequence 69, Appl
138	15	100.0	15	1	US-08-460-874A-26	Sequence 26, Appl	211	15	100.0	19	5	PCT-US94-01768-30	Sequence 30, Appl
139	15	100.0	15	1	US-08-133-271-2	Sequence 26, Appl	212	15	100.0	19	5	PCT-US95-03236-15	Sequence 15, Appl
140	15	100.0	15	1	US-08-133-271-3	Sequence 3, Appl	213	15	100.0	19	5	PCT-US95-03236-58	Sequence 58, Appl
141	15	100.0	15	2	US-08-967-101-167	Sequence 167, App	214	15	100.0	20	1	US-07-651-710A-17	Sequence 17, Appl
142	15	100.0	15	2	US-08-974-196-1	Sequence 1, Appl	215	15	100.0	20	1	US-08-080-809-3	Sequence 3, Appl
143	15	100.0	15	2	US-08-388-883B-26	Sequence 26, Appl	216	15	100.0	20	1	US-07-956-848A-24	Sequence 24, Appl
144	15	100.0	15	2	US-08-726-306A-60	Sequence 60, Appl	217	15	100.0	20	1	US-07-864-475A-5	Sequence 5, Appl
145	15	100.0	15	2	US-08-592-541-167	Sequence 167, App	218	15	100.0	20	1	US-08-460-874A-21	Sequence 21, Appl
146	15	100.0	15	3	US-09-124-698-167	Sequence 167, App	219	15	100.0	20	1	US-08-460-874A-23	Sequence 23, Appl
147	15	100.0	15	3	US-08-462-211A-26	Sequence 26, Appl	220	15	100.0	20	1	US-08-460-874A-24	Sequence 24, Appl
148	15	100.0	15	3	US-09-127-480-167	Sequence 167, App	221	15	100.0	20	1	US-08-471-956-24	Sequence 24, Appl
149	15	100.0	15	3	US-08-496-841C-164	Sequence 164, App	222	15	100.0	20	2	US-08-715-568A-13	Sequence 13, Appl
150	15	100.0	15	3	US-09-013-896A-29	Sequence 29, Appl	223	15	100.0	20	2	US-08-388-883B-23	Sequence 23, Appl
151	15	100.0	15	4	US-09-124-523-167	Sequence 167, App	224	15	100.0	20	2	US-08-388-883B-24	Sequence 24, Appl
152	15	100.0	15	4	US-09-636-796A-167	Sequence 167, App	225	15	100.0	20	2	US-08-468-249A-5	Sequence 5, Appl
153	15	100.0	15	4	US-09-827-948-29	Sequence 29, Appl	226	15	100.0	20	2	US-08-462-211A-21	Sequence 21, Appl
154	15	100.0	15	4	US-09-409-938-8	Sequence 8, Appl	227	15	100.0	20	3	US-08-462-211A-23	Sequence 23, Appl
155	15	100.0	15	4	US-08-685-852-2	Sequence 2, Appl	228	15	100.0	20	3	US-08-462-211A-24	Sequence 24, Appl
156	15	100.0	16	1	US-08-478-312-27	Sequence 27, Appl	229	15	100.0	20	3	US-08-817-895-7	Sequence 7, Appl
157	15	100.0	16	1	US-08-485-302-27	Sequence 27, Appl	230	15	100.0	20	3	US-08-817-895-8	Sequence 8, Appl
158	15	100.0	16	1	US-08-476-169-23	Sequence 23, Appl	231	15	100.0	20	4	US-09-144-428-64	Sequence 64, App
159	15	100.0	16	1	US-08-484-083-23	Sequence 23, Appl	232	15	100.0	20	4	US-09-428-082B-327	Sequence 327, App
160	15	100.0	16	2	US-08-373-190-26	Sequence 26, Appl	233	15	100.0	20	4	US-09-881-710-14	Sequence 14, Appl
161	15	100.0	16	2	US-08-902-623-6	Sequence 6, Appl	234	15	100.0	20	4		
162	15	100.0	16	2	US-08-438-190A-26	Sequence 26, Appl							
163	15	100.0	16	3	US-08-350-215-26	Sequence 26, Appl							
164	15	100.0	16	3	US-09-287-145A-26	Sequence 26, Appl							
165	15	100.0	16	3	US-08-564-164A-11	Sequence 11, Appl							
166	15	100.0	16	4	US-09-556-111-26	Sequence 11, Appl							
167	15	100.0	16	4	US-09-336-093-1	Sequence 1, Appl							
168	15	100.0	16	4	US-09-461-325-523	Sequence 523, App							
169	15	100.0	16	4	US-09-557-465D-1	Sequence 1, Appl							
170	15	100.0	16	4	US-10-012-542-523	Sequence 523, App							
171	15	100.0	16	5	PCT-US91-09422-22	Sequence 22, Appl							
172	15	100.0	17	1	US-08-168-809-21	Sequence 21, Appl							
173	15	100.0	17	1	US-08-071-811A-1	Sequence 1, Appl							

ALIGNMENTS

RESULT 1
US-08-864-301-1
; Sequence 1, Application US/08864301
; Patent No. 6126939
; GENERAL INFORMATION:
; APPLICANT: Eisenbach-Schwartz, M.
; APPLICANT: Beserman, P.

APPLICANT: Hirschberg, D.
TITLE OF INVENTION: ANTI-INFLAMMATORY PEPTIDES AND USES THEREOF
FILE REFERENCE: 5763-021
CURRENT APPLICATION NUMBER: US/08/864,301
CURRENT FILING DATE: 1997-05-28
EARLIER APPLICATION NUMBER: PCT/IL97/00295
EARLIER FILING DATE: 1997-09-03
EARLIER APPLICATION NUMBER: 08/864,301
EARLIER FILING DATE: 1997-05-28
EARLIER APPLICATION NUMBER: 08/753,141
EARLIER FILING DATE: 1996-11-20
EARLIER APPLICATION NUMBER: 60/025,376
EARLIER FILING DATE: 1996-09-03
NUMBER OF SEQ ID NOS: 1
SOFTWARE: Patent in Ver. 2.0
SEQ ID NO 1
LENGTH: 4
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: peptide
OTHER INFORMATION: derivative
US-08-864-301-1

Query Match 100.0%; Score 15; DB 3; Length 4;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 2 RER 4

RESULT 2
US-09-142-078-29
Sequence 29, Application US/09142078
Patent No. 6172041
GENERAL INFORMATION:
APPLICANT: McCabe, R. Tyler
APPLICANT: Zhou, Li-Ming
APPLICANT: Layer, Richard T.
TITLE OF INVENTION: Use of Conantokins
NUMBER OF SEQUENCES: 71
CORRESPONDENCE ADDRESS:
ADDRESSEE: Rothwell, Figg, Ernst & Kurz, P.C.
STREET: 555 Thirteenth Street, N.W., Suite 701-B
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent in Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/142,078
FILING DATE: 10-FEB-1999
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO US97/12652
FILING DATE: 21-JUL-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/762,377
FILING DATE: 06-DEC-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/684,750
FILING DATE: 22-JUL-1996
ATTORNEY/AGENT INFORMATION:
NAME: Ihnen, Jeffrey L.
REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 2314-135.A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-783-6040

TELEFAX: 202-783-6031
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
FEATURE:
NAME/KEY: Modified-site
LOCATION: 4
OTHER INFORMATION: /note= "Xaa is
OTHER INFORMATION: gamma-carboxyglutamic acid"
US-09-142-078-29

Query Match 100.0%; Score 15; DB 3; Length 4;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 1 RER 3

RESULT 3
US-09-357-141-29
Sequence 29, Application US/09357141
Patent No. 6277825
GENERAL INFORMATION:
APPLICANT: Olivera, Baldomero M.
APPLICANT: McIntosh, J. Michael
APPLICANT: McCabe, R. Tyler
APPLICANT: Layer, Richard T.
APPLICANT: Zhou, Li-Ming
TITLE OF INVENTION: Use of Conantokins for Treating Pain
FILE REFERENCE: 2314-171
CURRENT APPLICATION NUMBER: US/09/357,141
CURRENT FILING DATE: 1999-07-20
PRIOR APPLICATION NUMBER: US 09/283,277
PRIOR FILING DATE: 1999-04-01
PRIOR APPLICATION NUMBER: US 09/142,078
PRIOR FILING DATE: 1999-02-10
PRIOR APPLICATION NUMBER: WO US97/12652
PRIOR FILING DATE: 1997-07-21
PRIOR APPLICATION NUMBER: US 08/762,377
PRIOR FILING DATE: 1996-12-06
PRIOR APPLICATION NUMBER: US 08/684,750
PRIOR FILING DATE: 1996-07-22
NUMBER OF SEQ ID NOS: 71
SOFTWARE: Patent in Ver. 2.0
SEQ ID NO 29
LENGTH: 4
TYPE: PRT
ORGANISM: Conus sulcatus
FEATURE:
NAME/KEY: PEPTIDE
LOCATION: (4)
OTHER INFORMATION: Xaa is gamma-carboxyglutamic acid.
US-09-357-141-29

Query Match 100.0%; Score 15; DB 3; Length 4;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 1 RER 3

RESULT 4
US-09-533-889-29
Sequence 29, Application US/09533889

Patent No. 6399574
 GENERAL INFORMATION:
 APPLICANT: McCabe, R. Tyler
 APPLICANT: Zhou, Li-Ming
 APPLICANT: Laver, Richard T.
 APPLICANT: Olivera, Baldomero M.
 APPLICANT: McIntosh, J. Michael
 TITLE OF INVENTION: Use of Conantokins
 NUMBER OF SEQUENCES: 71
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Rothwell, Figg, Ernst & Kurz, P.C.
 STREET: 555 Thirteenth Street, N.W., Suite 701-E
 CITY: Washington
 STATE: D.C.
 COUNTRY: USA
 ZIP: 20004
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patent In Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/533,889
 FILING DATE: 22 MAR-2000
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 09/142,078
 FILING DATE: 10-FEB-1999
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: WO US97/12652
 FILING DATE: 21-JUL-1997
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/762,377
 FILING DATE: 06-DEC-1996
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/684,750
 FILING DATE: 22-JUL-1996
 ATTORNEY/AGENT INFORMATION:
 NAME: Ihnen, Jeffrey L.
 REGISTRATION NUMBER: 28,957
 REFERENCE/DOCKET NUMBER: 2314-168.A
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 202-783-6040
 TELEFAX: 202-783-6031
 INFORMATION FOR SEQ ID NO: 29:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 4 amino acids
 TYPE: amino acid
 STRANDEDNESS:
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 FRAGMENT TYPE: internal
 FEATURE:
 NAME/KEY: Modified-site
 LOCATION: 4
 OTHER INFORMATION: /note= "Xaa is
 gamma-carboxyglutamic acid"
 US-09-533-889-29

Query Match 100.0%; Score 15; DB 4; Length 4;
 Best Local Similarity 100.0%; Pred. No. 3e+05;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 Db 1 RER 3

RESULT 5
 US-09-142-080-29
 Sequence 29, Application US/09142080
 Patent No. 6515103
 GENERAL INFORMATION:
 APPLICANT: Abogadie, Fe C.

Cruz, Lourdes J.
 Olivera, Baldomero M.
 Walker, Craig
 Colledge, Clark
 Hillyard, David R.
 Jimenez, Elsie
 Laver, Richard T.
 Zhou, Li-Ming
 McCabe, R. Tyler
 TITLE OF INVENTION: Conantokins
 NUMBER OF SEQUENCES: 71
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Rothwell, Figg, Ernst & Manbeck, P.C.
 STREET: 555 Thirteenth Street, N.W., Suite 701-E
 CITY: Washington
 STATE: D.C.
 COUNTRY: USA
 ZIP: 20004
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patent In Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/142,080
 FILING DATE: 11-May-2000
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: WO US97/12618
 FILING DATE: 21-JUL-1997
 APPLICATION NUMBER: US 08/684,742
 FILING DATE: 22-JUL-1996
 ATTORNEY/AGENT INFORMATION:
 NAME: Ihnen, Jeffrey L.
 REGISTRATION NUMBER: 28,957
 REFERENCE/DOCKET NUMBER: 2314-134.A
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 202-783-6040
 TELEFAX: 202-783-6031
 INFORMATION FOR SEQ ID NO: 29:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 4 amino acids
 TYPE: amino acid
 STRANDEDNESS: <Unknown>
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 FRAGMENT TYPE: internal
 FEATURE:
 NAME/KEY: Modified-site
 LOCATION: 4
 OTHER INFORMATION: /note= "Xaa is
 gamma-carboxyglutamic acid"
 US-09-142-080-29

Query Match 100.0%; Score 15; DB 4; Length 4;
 Best Local Similarity 100.0%; Pred. No. 3e+05;
 Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
 Db 1 RER 3

RESULT 6
 PCT-US92-09070-14
 Sequence 14, Application PC/TUS9209070
 GENERAL INFORMATION:
 APPLICANT: Saich, Tsunao [NMI]
 TITLE OF INVENTION: SUBSTANCES HAVING THE GROWTH-PROMOTING
 EFFECT OF AMYLOID PRECURSOR PROTEIN
 NUMBER OF SEQUENCES: 18
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Knobbe, Martens, Olson and Bear

STREET: 620 Newport Center Drive
CITY: Newport Beach
STATE: California
COUNTRY: U.S.A.
ZIP: 92660
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/09070
FILING DATE: 19921023
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Altman, Daniel E
REGISTRATION NUMBER: 34115
REFERENCE/DOCKET NUMBER: UC035.001A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (714) 760-0404
TELEFAX: (714) 760-9502
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: internal
PCT-US92-09070-14

Query Match 100.0%; Score 15; DB 5; Length 4;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3

Db 1 RER 3

RESULT 7
US-08-552-142A-7
Sequence 7, Application US/08552142A
Patent No. 5695995
GENERAL INFORMATION:
APPLICANT: Weintraub, Harold M.
APPLICANT: Lee, Jacqueline E.
APPLICANT: Tapscoff, Stephen J.
APPLICANT: Hollenberg, Stanley M.
TITLE OF INVENTION: Neurogenic Differentiation (Neurod) Genes
NUMBER OF SEQUENCES: 20
TITLE OF INVENTION: and Proteins
CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor Johnson Kindness PLLC
STREET: 1420 Fifth Avenue, Suite 2800
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/552,142A
FILING DATE: 02-NOV-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/239,238
FILING DATE: 06-MAY-1994

PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US95/05741
FILING DATE: 08-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Broderick, Thomas F.
REGISTRATION NUMBER: 31,332
REFERENCE/DOCKET NUMBER: FHCR-1-8933
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-682-8100
TELEFAX: 206-225-0709
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
US-08-552-142A-7

Query Match 100.0%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3

Db 3 RER 5

RESULT 8
US-08-704-170-13
Sequence 13, Application US/08704170
Patent No. 5707626
GENERAL INFORMATION:
APPLICANT: Douvas, Angeline
APPLICANT: Takehana, Yoshi
APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 No. 5707626th Figueroa Street, Suite 500
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90012
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/704,170
FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-704-170-13

Query Match 100.0%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 1 RER 3

RESULT 9
US-08-704-170-14
; Sequence 14, Application US/08704170
; Patent No. 5707626
; GENERAL INFORMATION:
; APPLICANT: Douvas, Angeline
; APPLICANT: Takehana, Yoshi
; APPLICANT: Ehresmann, Glenn
; TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
; TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
; NUMBER OF SEQUENCES: 121
; CORRESPONDENCE ADDRESS: 121
; ADDRESSEE: Robbins, Berliner & Carson
; STREET: 201 No. 5707626th Figueroa Street, Suite 500
; CITY: Los Angeles
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 90012
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/704,170
; FILING DATE:
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/029,850
; FILING DATE: 11-MAR-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Spitals, John P.
; REGISTRATION NUMBER: 29,215
; REFERENCE/DOCKET NUMBER: 1920-331
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 977-1001
; TELEFAX: (213) 977-1003
; INFORMATION FOR SEQ ID NO: 14:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: Peptide
; US-08-704-170-14

Query Match 100.0%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 2 RER 4

RESULT 10
US-08-704-170-60
; Sequence 60, Application US/08704170
; Patent No. 5707626
; GENERAL INFORMATION:
; APPLICANT: Douvas, Angeline
; APPLICANT: Takehana, Yoshi
; APPLICANT: Ehresmann, Glenn
; TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
; TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS

; NUMBER OF SEQUENCES: 121
; CORRESPONDENCE ADDRESS: 121
; ADDRESSEE: Robbins, Berliner & Carson
; STREET: 201 No. 5707626th Figueroa Street, Suite 500
; CITY: Los Angeles
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 90012
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/704,170
; FILING DATE:
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/029,850
; FILING DATE: 11-MAR-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Spitals, John P.
; REGISTRATION NUMBER: 29,215
; REFERENCE/DOCKET NUMBER: 1920-331
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 977-1001
; TELEFAX: (213) 977-1003
; INFORMATION FOR SEQ ID NO: 60:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: Peptide
; US-08-704-170-60

Query Match 100.0%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 1 RER 3

RESULT 11
US-08-910-973-7
; Sequence 7, Application US/08910973
; Patent No. 5795723
; GENERAL INFORMATION:
; APPLICANT: Tapscott, Stephen J.
; APPLICANT: Olson, James M.
; TITLE OF INVENTION: Expression of Neurogenic bHLH Genes in Primitive Neuroectoder
; CORRESPONDENCE ADDRESS: 24
; ADDRESSEE: Christensen O'Connor Johnson Kindness PLLC
; STREET: 1420 Fifth Avenue, Suite 2800
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101-2347
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/910,973
; FILING DATE:
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/239,238
; FILING DATE: 06-MAY-1994
; PRIOR APPLICATION DATA:

APPLICATION NUMBER: WO PCT/US95/05741
FILING DATE: 08-MAY-1995
PRIORITY INFORMATION:
APPLICATION NUMBER: PCT/US96/17532
FILING DATE: 30-October-1996
ATTORNEY/AGENT INFORMATION:
NAME: Sheiness, Diana K.
REGISTRATION NUMBER: 35,356
REFERENCE/DOCKET NUMBER: FHCR-1-10958
TELEPHONE: 206-682-8100; 206-224-0735 (direct)
TELEFAX: 206-225-0779
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
US-09-910-973-7

Query Match 100.0%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 12
US-09-499-227-7
Sequence 7, Application US/09499227
Patent No. 644463
GENERAL INFORMATION:
APPLICANT: Tapscott, Stephen J.
TITLE OF INVENTION: Expression of Neurogenic bHLH Genes in Primitive Neuroectoderm
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor Johnson Kindness PLLC
STREET: 1420 Fifth Avenue, Suite 2800
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/499,227
FILING DATE: 05-August-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/239,238
FILING DATE: 06-May-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US95/05741
FILING DATE: 08-May-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/17532
FILING DATE: 30-October-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/910,973
FILING DATE: 07-August-1997
ATTORNEY/AGENT INFORMATION:
NAME: Sheiness, Diana K.
REGISTRATION NUMBER: 35,356
REFERENCE/DOCKET NUMBER: FHCR-1-12742
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-682-8100; 206-224-0735 (direct)
TELEFAX: 206-225-0779

INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
US-09-499-227-7

Query Match 100.0%; Score 15; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 13
US-09-638-202A-59
Sequence 59, Application US/09638202A
Patent No. 6462189
GENERAL INFORMATION:
APPLICANT: Koieda, Shohai
TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwesman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, Ste. 1600
CITY: Minneapolis
STATE: MN
COUNTRY: USA
ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/638,202A
FILING DATE: 11-Aug-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/096,749
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Ann S. Viksnins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
TELEFAX: (612) 339-3061
INFORMATION FOR SEQ ID NO: 59:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
SEQUENCE DESCRIPTION: SEQ ID NO: 59:
US-09-638-202A-59

Query Match 100.0%; Score 15; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 14
US-09-096-749A-59
; Sequence 59, Application US/09096749A
; Patent No. 6673901
; GENERAL INFORMATION:
; APPLICANT: Koieda, Shohei
; TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
; NUMBER OF SEQUENCES: 118
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Schwesman, Lundberg, Woessner & Kluth P.A.
; STREET: 121 South Eighth Street, Ste. 1600
; CITY: Minneapolis
; STATE: MN
; COUNTRY: USA
; ZIP: 55402
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 2.0b
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/096,749A
; FILING DATE: June 12, 1998
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; ATTORNEY/AGENT INFORMATION:
; NAME: Ann S. Vikenins
; REGISTRATION NUMBER: 37,748
; REFERENCE/DOCKET NUMBER: 109.034US1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (612) 373-6900
; TELEFAX: (612) 339-3061
; INFORMATION FOR SEQ ID NO: 59:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
US-09-096-749A-59
Query Match 100.0%; Score 15; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
Db 3 RER 5
RESULT 15
PCT-US92-09070-1
; Sequence 1, Application PC/TUS9209070
; GENERAL INFORMATION:
; APPLICANT: Saich, Tsunao [NNL]
; TITLE OF INVENTION: SUBSTANCES HAVING THE GROWTH-PROMOTING
; TITLE OF INVENTION: EFFECT OF AMYLOID PRECURSOR PROTEIN
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Knobbe, Martens, Olson and Bear
; STREET: 620 Newport Center Drive
; CITY: Newport Beach
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 92660
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US92/09070
; FILING DATE: 19921023
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Altman, Daniel E
; REGISTRATION NUMBER: 34,115
; REFERENCE/DOCKET NUMBER: UC035.001A
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (714) 760-0404
; TELEFAX: (714) 760-9502
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5 amino acids
; TYPE: AMINO ACID
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: internal
; PCT-US92-09070-1
Query Match 100.0%; Score 15; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
Db 1 RER 3
RESULT 16
PCT-US94-02631-13
; Sequence 13, Application PC/TUS9402631
; GENERAL INFORMATION:
; APPLICANT: Douvas, Angeline
; APPLICANT: Takehana, Yoshi
; APPLICANT: Ehresmann, Glenn
; TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
; TITLE OF INVENTION: IMMUNOINFECTION CLUSTER VIRUS INFECTIONS
; NUMBER OF SEQUENCES: 121
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Robbins, Berliner & Carson
; STREET: 201 North Figueroa Street, Suite 500
; CITY: Los Angeles
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 90012
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US94/02631
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/029,850
; FILING DATE: 11-MAR-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Spitals, John P.
; REGISTRATION NUMBER: 29,215
; REFERENCE/DOCKET NUMBER: 1920-331
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 977-1001
; TELEFAX: (213) 977-1003
; INFORMATION FOR SEQ ID NO: 13:
; SEQUENCE CHARACTERISTICS:

;
; LENGTH: 5 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
PCT-US94-02631-13

Query Match 100.0%; Score 15; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 1 RER 3

RESULT 17
PCT-US94-02631-14
; Sequence 14, Application PC/TUS9402631
; GENERAL INFORMATION:
; APPLICANT: Douvas, Angeline
; APPLICANT: Takehana, Yoshi
; APPLICANT: Ehrsman, Glenn
; TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
; IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
; NUMBER OF SEQUENCES: 121
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Robbins, Berliner & Carson
; STREET: 201 North Figueroa Street, Suite 500
; CITY: Los Angeles
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 90012
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US94/02631
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/029,850
; FILING DATE: 11-MAR-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Spitals, John P.
; REGISTRATION NUMBER: 29,215
; REFERENCE/DOCKET NUMBER: 1920-331
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 977-1001
; TELEFAX: (213) 977-1003
; INFORMATION FOR SEQ ID NO: 14:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
PCT-US94-02631-14

Query Match 100.0%; Score 15; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 2 RER 4

RESULT 18
PCT-US94-02631-60
; Sequence 60, Application PC/TUS9402631
; GENERAL INFORMATION:
; APPLICANT: Douvas, Angeline

;
; APPLICANT: Takehana, Yoshi
; APPLICANT: Ehrsman, Glenn
; TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
; IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
; NUMBER OF SEQUENCES: 121
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Robbins, Berliner & Carson
; STREET: 201 North Figueroa Street, Suite 500
; CITY: Los Angeles
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 90012
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US94/02631
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/029,850
; FILING DATE: 11-MAR-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Spitals, John P.
; REGISTRATION NUMBER: 29,215
; REFERENCE/DOCKET NUMBER: 1920-331
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 977-1001
; TELEFAX: (213) 977-1003
; INFORMATION FOR SEQ ID NO: 60:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
PCT-US94-02631-60

Query Match 100.0%; Score 15; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
DB 1 RER 3

RESULT 19
PCT-US95-05741-7
; Sequence 7, Application PC/TUS9505741
; GENERAL INFORMATION:
; APPLICANT: Weintraub, Harold
; APPLICANT: Lee, Jacqueline E.
; APPLICANT: Tapscott, Stephen J.
; APPLICANT: Hollenberg, Stanley M.
; TITLE OF INVENTION: Neurogenic Differentiation (Neurob) Gene
; NUMBER OF SEQUENCES: 11
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Christensen O'Connor Johnson Kindness
; STREET: 1420 Fifth Avenue, Suite 2800
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101-2347
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US95/05741

;; FILING DATE:
;; CLASSIFICATION:
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Broderick, Thomas F.
;; REGISTRATION NUMBER: 31,332
;; REFERENCE/DOCKET NUMBER: EPCR-1-8504
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: 206-682-8100
;; TELEFAX: 206-225-0709
;; INFORMATION FOR SEQ ID NO: 7:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 5 amino acids
;; TYPE: amino acid
;; TOPOLOGY: linear
;; MOLECULE TYPE: peptide
;; FRAGMENT TYPE: internal
PCT-US95-05741-7

Query Match 100.0%; Score 15; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
|||
Db 3 RER 5

RESULT 20
US-08-704-170-4
; Sequence 4, Application US/08704170
; Patent No. 5707626
; GENERAL INFORMATION:
; APPLICANT: Douvas, Angeline
; APPLICANT: Takehana, Yoshi
; APPLICANT: Ehresmann, Glenn
; TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
; TITLE OF INVENTION: IMMUNOINFECTION CLUSTER VIRUS INFECTIONS
; NUMBER OF SEQUENCES: 121
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Robbins, Berliner & Carson
; STREET: 201 No. 5707626th Figueroa Street, Suite 500
; CITY: Los Angeles
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 90012
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/704,170
; FILING DATE:
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/029,850
; FILING DATE: 11-MAR-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Spitals, John P.
; REGISTRATION NUMBER: 29,215
; REFERENCE/DOCKET NUMBER: 1920-331
; TELEPHONE: (213) 977-1001
; TELEFAX: (213) 977-1003
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 6 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-704-170-4

Query Match 100.0%; Score 15; DB 1; Length 6;

Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
|||
Db 2 RER 4

RESULT 21
US-08-704-170-97
; Sequence 97, Application US/08704170
; Patent No. 5707626
; GENERAL INFORMATION:
; APPLICANT: Douvas, Angeline
; APPLICANT: Takehana, Yoshi
; APPLICANT: Ehresmann, Glenn
; TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
; TITLE OF INVENTION: IMMUNOINFECTION CLUSTER VIRUS INFECTIONS
; NUMBER OF SEQUENCES: 121
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Robbins, Berliner & Carson
; STREET: 201 No. 5707626th Figueroa Street, Suite 500
; CITY: Los Angeles
; STATE: California
; COUNTRY: U.S.A.
; ZIP: 90012
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/704,170
; FILING DATE:
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/029,850
; FILING DATE: 11-MAR-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Spitals, John P.
; REGISTRATION NUMBER: 29,215
; REFERENCE/DOCKET NUMBER: 1920-331
; TELEPHONE: (213) 977-1001
; TELEFAX: (213) 977-1003
; INFORMATION FOR SEQ ID NO: 97:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 6 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-704-170-97

Query Match 100.0%; Score 15; DB 1; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
|||
Db 2 RER 4

RESULT 22
US-08-482-228-145
; Sequence 145, Application US/08482228
; Patent No. 5968753
; GENERAL INFORMATION:
; APPLICANT: Tseng-Law, Janet
; APPLICANT: Kobori, Joan A.
; APPLICANT: Al-Abdaly, Fahad A.
; APPLICANT: Guillermo, Roy
; APPLICANT: Helgeson, Sam L.
; APPLICANT: Deans, Robert J.

;; TITLE OF INVENTION: POSITIVE AND POSITIVE/NEGATIVE CELL
;; TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE RELEASE
;; NUMBER OF SEQUENCES: 215
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Janice Guthrie, Ph.D.
;; STREET: P.O. Box 15210
;; CITY: Irvine
;; STATE: California
;; COUNTRY: USA
;; ZIP: 92713-5210
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Floppy disk
;; COMPUTER: IBM PC compatible
;; OPERATING SYSTEM: PC-DOS/MS-DOS
;; SOFTWARE: PatentIn Release #1.0, Version #1.25
;; CURRENT APPLICATION DATA: US/08/482,228
;; APPLICATION NUMBER: US/08/482,228
;; FILING DATE: 07-JUN-1995
;; CLASSIFICATION: 435
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Guthrie, Janice
;; REGISTRATION NUMBER: 35,170
;; REFERENCE/DOCKET NUMBER: IT-4630CIP3
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (714) 440-5353
;; TELEFAX: (714) 553-1952
;; INFORMATION FOR SEQ ID NO: 145:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 6 amino acids
;; TYPE: amino acid
;; TOPOLOGY: linear
;; MOLECULE TYPE: peptide
US-08-482-228-145

Query Match 100.0%; Score 15; DB 2; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 1 RER 3

RESULT 23
US-08-482-528-145
; Sequence 145, Application US/08482528
; Patent No. 6017719
; GENERAL INFORMATION:
; APPLICANT: Tseng-Law, Janet
; APPLICANT: Kobori, Joan A.
; APPLICANT: Al-Abdaly, Fahad A.
; APPLICANT: Guillermo, Roy
; APPLICANT: Helgeson, Sam L.
; APPLICANT: Deans, Robert J.
; TITLE OF INVENTION: POSITIVE AND POSITIVE/NEGATIVE CELL
; TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE RELEASE
; NUMBER OF SEQUENCES: 215
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Janice Guthrie, Ph.D.
; STREET: P.O. Box 15210
; CITY: Irvine
; STATE: California
; COUNTRY: USA
; ZIP: 92713-5210
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/482,528
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 435

;; ATTORNEY/AGENT INFORMATION:
;; NAME: Guthrie, Janice
;; REGISTRATION NUMBER: 35,170
;; REFERENCE/DOCKET NUMBER: IT-4630CIP4
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (714) 440-5353
;; TELEFAX: (714) 553-1952
;; INFORMATION FOR SEQ ID NO: 145:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 6 amino acids
;; TYPE: amino acid
;; TOPOLOGY: linear
;; MOLECULE TYPE: peptide
US-08-482-528-145

Query Match 100.0%; Score 15; DB 3; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 1 RER 3

RESULT 24
US-09-020-880-60
; Sequence 60, Application US/09020880A
; Patent No. 6136558
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ballinger, Marcus D.
; APPLICANT: Jones, Jennifer T.
; APPLICANT: Fairbrother, Wayne J.
; APPLICANT: Slikowski, Mark X.
; APPLICANT: Wells, James A.
; TITLE OF INVENTION: HERGULIN VARIANTS
; FILE REFERENCE: 14918-720CON1
; CURRENT APPLICATION NUMBER: US/09/020,880A
; CURRENT FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: US 60/037,581
; EARLIER FILING DATE: 1997-02-10
; NUMBER OF SEQ ID NOS: 116
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 60
; LENGTH: 6
; TYPE: PRT
; ORGANISM: No. 6136558 relevant (recombinant)
US-09-020-880-60

Query Match 100.0%; Score 15; DB 3; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 2 RER 4

RESULT 25
US-09-101-544-60
; Sequence 60, Application US/09101544
; Patent No. 6387638
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ballinger, Marcus D.
; APPLICANT: Jones, Jennifer T.
; APPLICANT: Fairbrother, Wayne J.
; APPLICANT: Slikowski, Mark X.
; APPLICANT: Wells, James A.
; TITLE OF INVENTION: HERGULIN VARIANTS
; FILE REFERENCE: 14918-720CON2
; CURRENT APPLICATION NUMBER: US/09/101,544
; CURRENT FILING DATE: 1998-07-17

;; PRIOR APPLICATION NUMBER: US 09/020,880
;; PRIOR FILING DATE: 1998-02-09
;; PRIOR APPLICATION NUMBER: US 60/037,581
;; PRIOR FILING DATE: 1997-02-10
;; NUMBER OF SEQ ID NOS: 116
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 60
;; LENGTH: 6
;; TYPE: PRT
;; ORGANISM: No. 6387638 relevant (recombinant)
US-09-101-544-60

Query Match 100.0%; Score 15; DB 4; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 2 RER 4

RESULT 26

US-09-007-288E-21
;; Sequence 21, Application US/09007288E
;; Patent No. 6495357

GENERAL INFORMATION:

;; APPLICANT: Fugisang, Claus
;; APPLICANT: Okkels, Jens
;; APPLICANT: Petersen, Dorte
;; APPLICANT: Patkar, Shamkant
;; APPLICANT: Thellersen, Marianne
;; APPLICANT: Svendsen, Allan
;; APPLICANT: Borch, Kim
;; APPLICANT: Royer, John
;; APPLICANT: Kretzschmar, Titus
;; APPLICANT: Haikier, Torben
;; APPLICANT: Vind, Jesper
;; APPLICANT: Jorgensen, Steen

;; TITLE OF INVENTION: No. 6495357e1 Lipolytic Enzymes

;; FILE REFERENCE: 4455.404-US

;; CURRENT APPLICATION NUMBER: US/09/007,288E

;; CURRENT FILING DATE: 2000-01-14

;; NUMBER OF SEQ ID NOS: 162

;; SOFTWARE: PatentIn version 3.1

;; SEQ ID NO 21

;; LENGTH: 6

;; TYPE: PRT

;; ORGANISM: Artificial sequence

;; FEATURE:

;; OTHER INFORMATION: Peptide addition

US-09-007-288E-21

Query Match 100.0%; Score 15; DB 4; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 4 RER 6

RESULT 27

US-08-932-411A-29

;; Sequence 29, Application US/08932411A
;; Patent No. 6566496

GENERAL INFORMATION:

;; APPLICANT: Anderson, David J.
;; APPLICANT: Ma, Qiufu
;; TITLE OF INVENTION: NEUROGENIN
;; NUMBER OF SEQUENCES: 31
;; CORRESPONDENCE ADDRESS:

;; ADDRESSEE: Flehr Hobbach Test Albritton & Herbert LLP
;; STREET: Four Embarcadero Center, Suite 3400

;; CITY: San Francisco
;; STATE: California
;; COUNTRY: United States
;; ZIP: 94111-4187
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Floppy disk
;; COMPUTER: IBM PC compatible
;; OPERATING SYSTEM: PC-DOS/MS-DOS
;; SOFTWARE: PatentIn Release #1.0, Version #1.30
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/08/932,411A
;; FILING DATE: 15-SEP-1997
;; CLASSIFICATION: 536

;; PRIOR APPLICATION DATA:

;; APPLICATION NUMBER: US 08/772,009

;; FILING DATE: 19-DEC-1996

;; PRIOR APPLICATION DATA:

;; APPLICATION NUMBER: US 08/722,570

;; FILING DATE: 19-DEC-1996

;; ATTORNEY/AGENT INFORMATION:

;; NAME: Silva, Robin W.

;; REGISTRATION NUMBER: 38,304

;; REFERENCE/DOCKET NUMBER: A-63902-3/RPT/RMS

;; TELECOMMUNICATION INFORMATION:

;; TELEPHONE: (415) 781-1989

;; TELEFAX: (415) 398-3249

;; TELEX: 910 277299

;; INFORMATION FOR SEQ ID NO: 29:

;; SEQUENCE CHARACTERISTICS:

;; LENGTH: 6 amino acids

;; TYPE: amino acid

;; STRANDEDNESS: unknown

;; TOPOLOGY: unknown

;; MOLECULE TYPE: protein

US-08-932-411A-29

Query Match 100.0%; Score 15; DB 4; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 4 RER 6

RESULT 28

US-08-932-411A-30

;; Sequence 30, Application US/08932411A
;; Patent No. 6566496

GENERAL INFORMATION:

;; APPLICANT: Anderson, David J.

;; APPLICANT: Ma, Qiufu

;; TITLE OF INVENTION: NEUROGENIN

;; NUMBER OF SEQUENCES: 31

;; CORRESPONDENCE ADDRESS:

;; ADDRESSEE: Flehr Hobbach Test Albritton & Herbert LLP
;; STREET: Four Embarcadero Center, Suite 3400

;; CITY: San Francisco

;; STATE: California

;; COUNTRY: United States

;; ZIP: 94111-4187

;; COMPUTER READABLE FORM:

;; MEDIUM TYPE: Floppy disk

;; COMPUTER: IBM PC compatible

;; OPERATING SYSTEM: PC-DOS/MS-DOS

;; SOFTWARE: PatentIn Release #1.0, Version #1.30

;; CURRENT APPLICATION DATA:

;; APPLICATION NUMBER: US/08/932,411A

;; FILING DATE: 15-SEP-1997

;; CLASSIFICATION: 536

;; PRIOR APPLICATION DATA:

;; APPLICATION NUMBER: US 08/772,009

;; FILING DATE: 19-DEC-1996

;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: US 08/722,570
;; FILING DATE: 19-DEC-1996
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Silva, Robin M.
;; REGISTRATION NUMBER: 38,304
;; REFERENCE/DOCKET NUMBER: A-63902-3/RFT/RMS
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (415) 781-1989
;; TELEFAX: (415) 398-3249
;; TELEX: 910 277299
;; INFORMATION FOR SEQ ID NO: 30:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 6 amino acids
;; TYPE: amino acid
;; STRANDEDNESS: unknown
;; TOPOLOGY: unknown
;; MOLECULE TYPE: protein
;; US-08-932-411A-30

Query Match 100.0%; Score 15; DB 4; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 4 RER 6

RESULT 29
PCT-US92-09070-13
;; Sequence 13, Application PC/TUS9209070
;; GENERAL INFORMATION:
;; APPLICANT: Saitoh, Tsunao [NM1]
;; TITLE OF INVENTION: SUBSTANCES HAVING THE GROWTH-PROMOTING
;; TITLE OF INVENTION: EFFECT OF AMYLOID PRECURSOR PROTEIN
;; NUMBER OF SEQUENCES: 18
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Knobbe, Martens, Olson and Bear
;; STREET: 620 Newport Center Drive
;; CITY: Newport Beach
;; STATE: California
;; COUNTRY: U.S.A.
;; ZIP: 92660
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Floppy disk
;; COMPUTER: IBM PC compatible
;; OPERATING SYSTEM: PC-DOS/MS-DOS
;; SOFTWARE: Patent In Release #1.0, Version #1.25
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: PCT/US92/09070
;; FILING DATE: 19921023
;; CLASSIFICATION: 514
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Altman, Daniel E
;; REGISTRATION NUMBER: 34,115
;; REFERENCE/DOCKET NUMBER: UC035.001A
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (714) 760-0404
;; TELEFAX: (714) 760-9502
;; INFORMATION FOR SEQ ID NO: 13:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 6 amino acids
;; TYPE: AMINO ACID
;; STRANDEDNESS: single
;; TOPOLOGY: linear
;; MOLECULE TYPE: peptide
;; HYPOTHETICAL: NO
;; ANTI-SENSE: NO
;; FRAGMENT TYPE: internal
;; PCT-US92-09070-13

Query Match 100.0%; Score 15; DB 5; Length 6;

Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 2 RER 4

RESULT 30
PCT-US94-02631-4
;; Sequence 4, Application PC/TUS9402631
;; GENERAL INFORMATION:
;; APPLICANT: Douvas, Angeline
;; APPLICANT: Takehana, Yoshi
;; APPLICANT: Ehresmann, Glenn
;; TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
;; TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
;; NUMBER OF SEQUENCES: 121
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Robbins, Berliner & Carson
;; STREET: 201 North Figueroa Street, Suite 500
;; CITY: Los Angeles
;; STATE: California
;; COUNTRY: U.S.A.
;; ZIP: 90012
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Floppy disk
;; COMPUTER: IBM PC compatible
;; OPERATING SYSTEM: PC-DOS/MS-DOS
;; SOFTWARE: Patent In Release #1.0, Version #1.25
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: PCT/US94/02631
;; FILING DATE:
;; CLASSIFICATION:
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: US 08/029,850
;; FILING DATE: 11-MAR-1993
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Spitals, John P.
;; REGISTRATION NUMBER: 29,215
;; REFERENCE/DOCKET NUMBER: 1920-331
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (213) 977-1001
;; TELEFAX: (213) 977-1003
;; INFORMATION FOR SEQ ID NO: 4:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 6 amino acids
;; TYPE: amino acid
;; TOPOLOGY: linear
;; MOLECULE TYPE: peptide
;; PCT-US94-02631-4

Query Match 100.0%; Score 15; DB 5; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 2 RER 4

RESULT 31
PCT-US94-02631-97
;; Sequence 97, Application PC/TUS9402631
;; GENERAL INFORMATION:
;; APPLICANT: Douvas, Angeline
;; APPLICANT: Takehana, Yoshi
;; APPLICANT: Ehresmann, Glenn
;; TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
;; TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
;; NUMBER OF SEQUENCES: 121
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Robbins, Berliner & Carson

STREET: 201 North Figueroa Street, Suite 500
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90012
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02631
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 1920-331
TELEPHONE: (213) 977-1001
TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 97:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
PCT-US94-02631-97

Query Match 100.0%; Score 15; DB 5; Length 6;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
DB 2 RER 4

RESULT 32
US-08-552-142A-5
Sequence 5, Application US/08552142A
Patent No. 5695995
GENERAL INFORMATION:
APPLICANT: Weintraub, Harold M.
APPLICANT: Lee, Jacqueline E.
APPLICANT: Tapscott, Stephen J.
APPLICANT: Hollenberg, Stanley M.
TITLE OF INVENTION: Neurogenic Differentiation (Neurod) Genes
TITLE OF INVENTION: and Proteins
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor Johnson Kindness PLLC
STREET: 1420 Fifth Avenue, Suite 2800
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/552,142A
FILING DATE: 02-NOV-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/239,238
FILING DATE: 06-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US95/05741
FILING DATE: 08-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Broderick, Thomas F.
REGISTRATION NUMBER: 31,332
REFERENCE/DOCKET NUMBER: FHC-1-8933
TELEPHONE: 206-682-8100
TELEFAX: 206-225-0709
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
TYPE: amino acid
TOPOLOGY: linear

APPLICATION NUMBER: WO PCT/US95/05741
FILING DATE: 08-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Broderick, Thomas F.
REGISTRATION NUMBER: 31,332
REFERENCE/DOCKET NUMBER: FHC-1-8933
TELEPHONE: 206-682-8100
TELEFAX: 206-225-0709
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
US-08-552-142A-5

Query Match 100.0%; Score 15; DB 1; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
DB 3 RER 5

RESULT 33
US-08-552-142A-6
Sequence 6, Application US/08552142A
Patent No. 5695995
GENERAL INFORMATION:
APPLICANT: Weintraub, Harold M.
APPLICANT: Lee, Jacqueline E.
APPLICANT: Tapscott, Stephen J.
APPLICANT: Hollenberg, Stanley M.
TITLE OF INVENTION: Neurogenic Differentiation (Neurod) Genes
TITLE OF INVENTION: and Proteins
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor Johnson Kindness PLLC
STREET: 1420 Fifth Avenue, Suite 2800
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/552,142A
FILING DATE: 02-NOV-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/239,238
FILING DATE: 06-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US95/05741
FILING DATE: 08-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Broderick, Thomas F.
REGISTRATION NUMBER: 31,332
REFERENCE/DOCKET NUMBER: FHC-1-8933
TELEPHONE: 206-682-8100
TELEFAX: 206-225-0709
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
TYPE: amino acid
TOPOLOGY: linear

; MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-552-142A-6

Query Match 100.0%; Score 15; DB 1; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 3 RER 5

RESULT 34
US-08-910-973-5
; Sequence 5, Application US/08910973
; Patent No. 5795723
; GENERAL INFORMATION:
; APPLICANT: Tapscott, Stephen J.
; APPLICANT: Olson, James M.
; TITLE OF INVENTION: Expression of Neurogenic bHLH Genes in Primitive Neuroectoder
; NUMBER OF SEQUENCES: 24
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Christensen O'Connor Johnson KindnessPLLC
; STREET: 1420 Fifth Avenue, Suite 2800
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101-2347
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/910,973
; FILING DATE:
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/239,238
; FILING DATE: 06-MAY-1994
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: WO PCT/US95/05741
; FILING DATE: 08-MAY-1995
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: PCT/US96/17532
; FILING DATE: 30-October-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Sheiness, Diana K.
; REGISTRATION NUMBER: 35,356
; REFERENCE/DOCKET NUMBER: FHCR-1-10958
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-682-8100; 206-224-0735 (direct)
; TELEFAX: 206-225-0779
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 7 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-910-973-5

Query Match 100.0%; Score 15; DB 1; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 3 RER 5

RESULT 35

US-08-910-973-6
; Sequence 6, Application US/08910973
; Patent No. 5795723
; GENERAL INFORMATION:
; APPLICANT: Tapscott, Stephen J.
; APPLICANT: Olson, James M.
; TITLE OF INVENTION: Expression of Neurogenic bHLH Genes in Primitive Neuroectoder
; NUMBER OF SEQUENCES: 24
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Christensen O'Connor Johnson KindnessPLLC
; STREET: 1420 Fifth Avenue, Suite 2800
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101-2347
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/910,973
; FILING DATE:
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/239,238
; FILING DATE: 06-MAY-1994
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: WO PCT/US95/05741
; FILING DATE: 08-MAY-1995
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: PCT/US96/17532
; FILING DATE: 30-October-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Sheiness, Diana K.
; REGISTRATION NUMBER: 35,356
; REFERENCE/DOCKET NUMBER: FHCR-1-10958
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-682-8100; 206-224-0735 (direct)
; TELEFAX: 206-225-0779
; INFORMATION FOR SEQ ID NO: 6:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 7 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; FRAGMENT TYPE: internal
US-08-910-973-6

Query Match 100.0%; Score 15; DB 1; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
Db 3 RER 5

RESULT 36
US-08-907-403A-5
; Sequence 5, Application US/08907403A
; Patent No. 6013633
; GENERAL INFORMATION:
; APPLICANT: Balasubramaniam, Ambikaipakan
; APPLICANT: Chance, William T.
; TITLE OF INVENTION: Compounds For Control
; TITLE OF INVENTION: Of Appetite, Blood Pressure, Cardiovascular
; TITLE OF INVENTION: Response, Libido, And Circadian Rhythm
; NUMBER OF SEQUENCES: 8
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Wood, Herron & Evans, L.L.P.
; STREET: 441 Vine Street
; CITY: Cincinnati

STATE: Ohio
COUNTRY: USA
ZIP: 45202-2917
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
1.44 MB storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: Microsoft Word
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/907,403A
FILING DATE: 07-AUG-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/023,588
FILING DATE: 09-AUG-1996
ATTORNEY/AGENT INFORMATION:
NAME: Albainy-Jenei, Stephen R.
REGISTRATION NUMBER: 41,487
REFERENCE/DOCKET NUMBER: UOC-113A-111
TELECOMMUNICATION INFORMATION:
TELEPHONE: (513) 241-2324
TELEFAX: (513) 421-7269
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 7
TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
ANTI-SENSE: no
FEATURE:
LOCATION: 1
OTHER INFORMATION: Xaa represents Asp
US-08-907-403A-5
Query Match 100.0%; Score 15; DB 3; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
DB 4 RER 6
RESULT 37
US-08-907-403A-6
Sequence 6, Application US/08907403A
Patent No. 6013633
GENERAL INFORMATION:
APPLICANT: Balasubramaniam, Ambikaipakan
APPLICANT: Chance, William T.
TITLE OF INVENTION: Compounds For Control
TITLE OF INVENTION: Of Appetite, Blood Pressure, Cardiovascular
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wood, Heron & Evans, L.L.P.
STREET: 441 Vine Street
CITY: Cincinnati
STATE: Ohio
COUNTRY: USA
ZIP: 45202-2917
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
1.44 MB storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: Microsoft Word
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/907,403A
FILING DATE: 07-AUG-1997

CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/023,588
FILING DATE: 09-AUG-1996
ATTORNEY/AGENT INFORMATION:
NAME: Albainy-Jenei, Stephen R.
REGISTRATION NUMBER: 41,487
REFERENCE/DOCKET NUMBER: UOC-113A-111
TELECOMMUNICATION INFORMATION:
TELEPHONE: (513) 241-2324
TELEFAX: (513) 421-7269
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 7
TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: no
ANTI-SENSE: no
FEATURE:
LOCATION: 1
OTHER INFORMATION: Xaa represents Asp
US-08-907-403A-6
Query Match 100.0%; Score 15; DB 3; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
DB 4 RER 6
RESULT 38
US-09-499-227-5
Sequence 5, Application US/09499227
Patent No. 6444663
GENERAL INFORMATION:
APPLICANT: Tapscott, Stephen J.
APPLICANT: Olson, James M.
TITLE OF INVENTION: Expression of Neurogenic bHLH Genes in Primitive Neuroectoder
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor Johnson Kindness PLLC
STREET: 1420 Fifth Avenue, Suite 2800
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/499,227
FILING DATE: 05-August-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/239,238
FILING DATE: 06-May-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US95/05741
FILING DATE: 08-May-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/17532
FILING DATE: 30-October-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/910,973
FILING DATE: 07-August-1997
ATTORNEY/AGENT INFORMATION:
NAME: Sheiness, Diana K.
REGISTRATION NUMBER: 35,356
REFERENCE/DOCKET NUMBER: FHCR-1-12742
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-682-8100; 206-224-0735 (direct)

TELEFAX: 206-225-0779
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
US-09-499-227-5

Query Match 100.0%; Score 15; DB 4; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
DB 3 RER 5

RESULT 39

US-09-499-227-6
Sequence 6, Application US/09499227
Patent No. 644463
GENERAL INFORMATION:
APPLICANT: Tapscott, Stephen J.
APPLICANT: Olson, James M.
TITLE OF INVENTION: Expression of Neurogenic bHLH Genes in Primitive Neuroectoderm
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor Johnson Kindness PLLC
STREET: 1420 Fifth Avenue, Suite 2800
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/499,227
FILING DATE: 05-August-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/239,238
FILING DATE: 06-May-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US95/05741
FILING DATE: 08-May-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/17532
FILING DATE: 30-October-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/910,973
FILING DATE: 07-August-1997
ATTORNEY/AGENT INFORMATION:
NAME: Sheiness, Diana K.
REGISTRATION NUMBER: 35,356
REFERENCE/DOCKET NUMBER: PFCR-1-12742
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-682-8100; 206-224-0735 (direct)
TELEFAX: 206-225-0779
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
US-09-499-227-6

Query Match 100.0%; Score 15; DB 4; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;

Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 RER 3
|||
DB 3 RER 5

RESULT 40

US-09-007-288E-49
Sequence 49, Application US/09007288E
Patent No. 6495357
GENERAL INFORMATION:
APPLICANT: Fuglsang, Claus
APPLICANT: Okkels, Jens
APPLICANT: Petersen, Dorte
APPLICANT: Patkar, Shamkant
APPLICANT: Thellersen, Marianne
APPLICANT: Svenden, Allan
APPLICANT: Borch, Kim
APPLICANT: Royer, John
APPLICANT: Kretzschmar, Ritus
APPLICANT: Halkier, Torben
APPLICANT: Vind, Jesper
APPLICANT: Jorgensen, Steen
TITLE OF INVENTION: No. 6495357e1 Lipolytic Enzymes
FILE REFERENCE: 4455.404-US
CURRENT APPLICATION NUMBER: US/09/007,288E
CURRENT FILING DATE: 2000-01-14
NUMBER OF SEQ ID NOS: 162
SOFTWARE: Patent In version 3.1
SEQ ID NO 49
LENGTH: 7
TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
OTHER INFORMATION: Peptide addition
US-09-007-288E-49

Query Match 100.0%; Score 15; DB 4; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
DB 4 RER 6

RESULT 41

PCT-US95-05741-5
Sequence 5, Application PC/TUS9505741
GENERAL INFORMATION:
APPLICANT: Weintraub, Harold
APPLICANT: Lee, Jacqueline E.
APPLICANT: Tapscott, Stephen J.
APPLICANT: Hollenberg, Stanley M.
TITLE OF INVENTION: Neurogenic Differentiation (NeuroD) Gene
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor Johnson Kindness
STREET: 1420 Fifth Avenue, Suite 2800
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/05741
FILING DATE:

CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Broderick, Thomas F.
REGISTRATION NUMBER: 31,332
REFERENCE/DOCKET NUMBER: FHCR-1-8504
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-682-8100
TELEFAX: 206-225-0709
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
PCT-US95-05741-5

Query Match 100.0%; Score 15; DB 5; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
DB 3 RER 5

RESULT 42
PCT-US95-05741-6
Sequence 6, Application PC/TUS9505741
GENERAL INFORMATION:
APPLICANT: Weintraub, Harold
APPLICANT: Lee, Jacqueline E.
APPLICANT: Tapscoott, Stephen J.
APPLICANT: Hollenberg, Stanley M.
TITLE OF INVENTION: Neurogenic Differentiation (Neurod) Gene
TITLE OF INVENTION: and Protein
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor Johnson Kindness
STREET: 1420 Fifth Avenue, Suite 2800
CITY: Seattle
STATE: WA
COUNTRY: USA
ZIP: 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/05741
FILING DATE:

CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Broderick, Thomas F.
REGISTRATION NUMBER: 31,332
REFERENCE/DOCKET NUMBER: FHCR-1-8504
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-682-8100
TELEFAX: 206-225-0709
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
PCT-US95-05741-6
Query Match 100.0%; Score 15; DB 5; Length 7;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
DB 3 RER 5

RESULT 43
US-08-259-550A-35
Sequence 35, Application US/08259550A
Patent No. 5778892
GENERAL INFORMATION:
APPLICANT: Counts, David F.
APPLICANT: Duff, Ronald G.
TITLE OF INVENTION: Anti-Inflammatory Peptides
NUMBER OF SEQUENCES: 91
CORRESPONDENCE ADDRESS:
ADDRESSEE: PENNIE & EDMONDS
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/259,550A
FILING DATE: 16-JUN-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Miarock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 7142-011
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
TOPOLOGY: unknown
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-259-550A-35

Query Match 100.0%; Score 15; DB 1; Length 8;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
DB 6 RER 8

RESULT 44
US-08-461-216-7
Sequence 7, Application US/08461216
Patent No. 5958883
GENERAL INFORMATION:
APPLICANT: Snow, A.D.
TITLE OF INVENTION: ANIMAL MODELS OF HUMAN AMYLOIDOSES
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen, O'Connor, Johnson and Kindness
STREET: 1420 Fifth Avenue, Suite 2800
CITY: Seattle
STATE: Washington
COUNTRY: USA
ZIP: 98101-2347
COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette-5.25 inch, 1.2Mb storage
COMPUTER: IBM PC/386 Compatible
OPERATING SYSTEM: MS-DOS 4.01
SOFTWARE: Word for Windows-t
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,216
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,734
FILING DATE: October 23, 1992
APPLICATION NUMBER: 07/950,417
FILING DATE: September 23, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Broderick, Thomas F.
REGISTRATION NUMBER: 31,332
REFERENCE/DOCKET NUMBER: UOPW-1-6707
TELECOMMUNICATION INFORMATION:
TELEPHONE: 1-206-682-8100; 1-206-224-0709 (direct)
TELEFAX: 1-206-224-0779
TELEX: 4938023
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
DESCRIPTION: APP(324-331); page 84, lines 8-13
US-08-461-216-7

Query Match 100.0%; Score 15; DB 2; Length 8;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 5 RER 7

RESULT 45
US-09-105-839D-25
; Sequence 25, Application US/09105839D
; Patent No. 6287756
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Chen, Yao-Tseng
; APPLICANT: Sahin, Ugur
; APPLICANT: Gure, Ali
; APPLICANT: Old, Lloyd J
; APPLICANT: Pfreundschuh, Michael
; TITLE OF INVENTION: Method for Determining Presence of Cancer In A Sample By Determin
; TITLE OF INVENTION: Expression of an SSX gene
; FILE REFERENCE: LUD 5556
; CURRENT APPLICATION NUMBER: US/09/105,839D
; CURRENT FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 72
; SEQ ID NO 25
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-105-839D-25

Query Match 100.0%; Score 15; DB 3; Length 8;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 46
US-09-105-839D-40
; Sequence 40, Application US/09105839D
; Patent No. 6287756
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Chen, Yao-Tseng
; APPLICANT: Sahin, Ugur
; APPLICANT: Gure, Ali
; APPLICANT: Old, Lloyd J
; APPLICANT: Pfreundschuh, Michael
; TITLE OF INVENTION: Method for Determining Presence of Cancer In A Sample By Determin
; TITLE OF INVENTION: Expression of an SSX gene
; FILE REFERENCE: LUD 5556
; CURRENT APPLICATION NUMBER: US/09/105,839D
; CURRENT FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 72
; SEQ ID NO 40
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-105-839D-40

Query Match 100.0%; Score 15; DB 3; Length 8;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 47
US-09-105-839D-57
; Sequence 57, Application US/09105839D
; Patent No. 6287756
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Chen, Yao-Tseng
; APPLICANT: Sahin, Ugur
; APPLICANT: Gure, Ali
; APPLICANT: Old, Lloyd J
; APPLICANT: Pfreundschuh, Michael
; TITLE OF INVENTION: Method for Determining Presence of Cancer In A Sample By Determin
; TITLE OF INVENTION: Expression of an SSX gene
; FILE REFERENCE: LUD 5556
; CURRENT APPLICATION NUMBER: US/09/105,839D
; CURRENT FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 72
; SEQ ID NO 57
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-105-839D-57

Query Match 100.0%; Score 15; DB 3; Length 8;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 RER 3
|||
Db 3 RER 5

RESULT 48
US-09-105-839D-69
; Sequence 69, Application US/09105839D
; Patent No. 6287756


```
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Chen, Yao-Tseng
; APPLICANT: Sahin, Ugur
; APPLICANT: Gure, Ali
; APPLICANT: Old, Lloyd J
; APPLICANT: Pfrendschuh, Michael
; TITLE OF INVENTION: Method for Determining Presence of Cancer In A Sample By Determining
; FILE REFERENCE: LUD 5556
; CURRENT APPLICATION NUMBER: US/09/105,839D
; CURRENT FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 72
; SEQ ID NO 69
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-105-839D-69

Query Match 100.0%; Score 15; DB 3; Length 8;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 3 RER 5

RESULT 49
US-08-723-661B-7
; Sequence 7, Application US/08723661B
; Patent No. 6340783
; GENERAL INFORMATION:
; APPLICANT: Alan D Snow
; TITLE OF INVENTION: Animal Models of Human Amyloidoses
; NUMBER OF SEQUENCES: 7
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Patrick M. Dwyer
; STREET: 1818 Westlake Avenue N, Suite 114
; CITY: Seattle
; STATE: WA (Washington)
; COUNTRY: United States of America
; ZIP: 98109
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette - 3.50 inch, 1.44 Mb storage
; COMPUTER: IBM PC
; OPERATING SYSTEM: PC-DOS (Windows 98)
; SOFTWARE: WordPerfect 5.2
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/723,661B
; FILING DATE: 31-Oct-1996
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/461,216
; FILING DATE: 05-Jun-1995
; APPLICATION NUMBER: 07/969,734
; FILING DATE: 23-Oct-1992
; APPLICATION NUMBER: 07/950,417
; FILING DATE: 23-Sep-1992
; ATTORNEY/AGENT INFORMATION:
; NAME: Dwyer, Patrick M.
; REGISTRATION NUMBER: 32,411
; REFERENCE/DOCKET NUMBER: PROTEO.P00C1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (206) 343-7074
; TELEFAX: (206) 343-7085
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 8 AMINO ACIDS
; TYPE: AMINO ACID
; STRANDEDNESS: SINGLE
; TOPOLOGY: LINEAR
;
```

```
; MOLECULE TYPE: PEPTIDE
; DESCRIPTION: APP (324-331); page 84, lines 8-13
; SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-08-723-661B-7

Query Match 100.0%; Score 15; DB 4; Length 8;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 5 RER 7

RESULT 50
US-09-344-040C-39
; Sequence 39, Application US/09344040C
; Patent No. 6548064
; GENERAL INFORMATION:
; APPLICANT: Tureci, Ozlem
; APPLICANT: Sahin, Ugur
; APPLICANT: Pfrendschuh, Michael
; APPLICANT: Ramensee, Hans Georg
; APPLICANT: Stevanovic, Stefan
; TITLE OF INVENTION: Method for Determining Presence of Cancer In a Sample By Determining
; TITLE OF INVENTION: Expression of an SSX Gene, Peptides Derived From Said SSX Gene ar
; TITLE OF INVENTION: Gene, and Uses Thereof
; FILE REFERENCE: LUD 5556.1
; CURRENT APPLICATION NUMBER: US/09/344,040C
; CURRENT FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 09/105,839
; PRIOR FILING DATE: 1998-06-26
; PRIOR APPLICATION NUMBER: US 08/851,130
; PRIOR FILING DATE: 1997-05-05
; NUMBER OF SEQ ID NOS: 132
; SEQ ID NO 39
; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-344-040C-39

Query Match 100.0%; Score 15; DB 4; Length 8;
Best Local Similarity 100.0%; Pred. No. 3e+05;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 RER 3
Db 3 RER 5

Search completed: March 5, 2004, 16:11:42
Job time : 25 secs
```

=> fil reg; d stat que 15
FILE 'REGISTRY' ENTERED AT 12:04:31 ON 08 MAR 2004
USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.
PLEASE SEE "HELP USAGETERMS" FOR DETAILS.
COPYRIGHT (C) 2004 American Chemical Society (ACS)

Property values tagged with IC are from the ZIC/VINITI data file
provided by InfoChem.

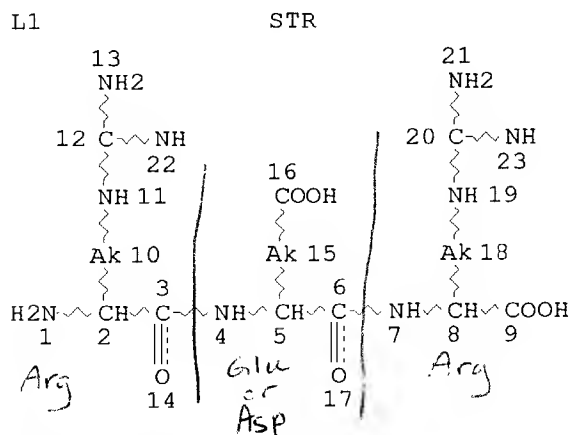
STRUCTURE FILE UPDATES: 7 MAR 2004 HIGHEST RN 659718-58-8
DICTIONARY FILE UPDATES: 7 MAR 2004 HIGHEST RN 659718-58-8

TSCA INFORMATION NOW CURRENT THROUGH JANUARY 6, 2004

Please note that search-term pricing does apply when
conducting SmartSELECT searches.

Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. For more
information enter HELP PROP at an arrow prompt in the file or refer
to the file summary sheet on the web at:
<http://www.cas.org/ONLINE/DBSS/registryss.html>



NODE ATTRIBUTES:
CONNECT IS E2 RC AT 10
CONNECT IS E2 RC AT 15
CONNECT IS E2 RC AT 18
DEFAULT MLEVEL IS ATOM
DEFAULT ECLEVEL IS LIMITED

GRAPH ATTRIBUTES:
RING(S) ARE ISOLATED OR EMBEDDED
NUMBER OF NODES IS 23

STEREO ATTRIBUTES: NONE
L5 2 SEA FILE=REGISTRY SSS FUL L1

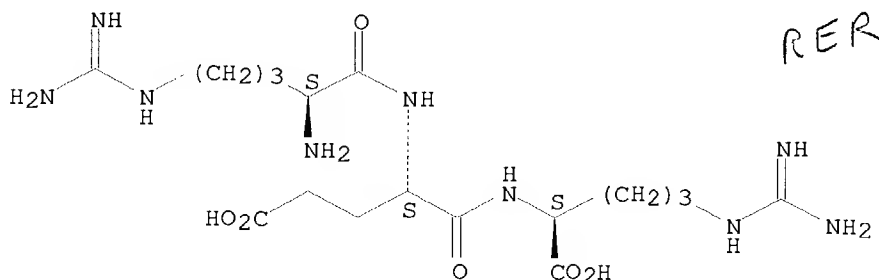
100.0% PROCESSED 66592 ITERATIONS
SEARCH TIME: 00.00.17

2 ANSWERS

=> d sqide 15 1-2

L5 ANSWER 1 OF 2 REGISTRY COPYRIGHT 2004 ACS on STN
RN 148914-10-7 REGISTRY *Use Registry # to match structure to citation*
CN L-Arginine, L-arginyl-L-.alpha.-glutamyl- (9CI) (CA INDEX NAME)
OTHER CA INDEX NAMES:
CN L-Arginine, N2-(N-L-arginyl-L-.alpha.-glutamyl)-
OTHER NAMES:
CN 9: PN: WO02083729 SEQID: 9 claimed sequence
FS STEREOSEARCH
MF C17 H33 N9 O6
SR CA
LC STN Files: CA, CAPLUS, TOXCENTER, USPATFULL

Absolute stereochemistry.

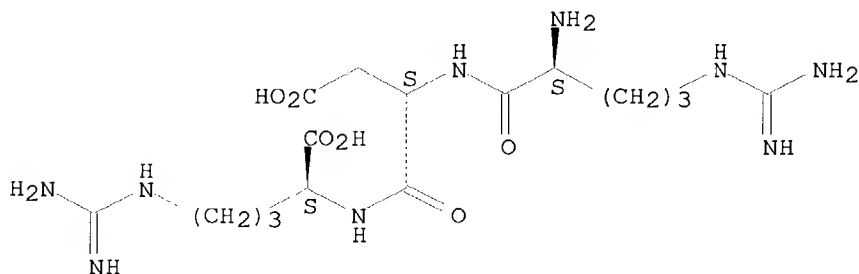


PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT

5 REFERENCES IN FILE CA (1907 TO DATE)
5 REFERENCES IN FILE CAPLUS (1907 TO DATE)

L5 ANSWER 2 OF 2 REGISTRY COPYRIGHT 2004 ACS on STN
RN 106326-11-8 REGISTRY
CN L-Arginine, N2-(N-L-arginyl-L-.alpha.-aspartyl)- (9CI) (CA INDEX NAME)
FS STEREOSEARCH
MF C16 H31 N9 O6
SR CA
LC STN Files: CA, CAPLUS, TOXCENTER

Absolute stereochemistry.



PROPERTY DATA AVAILABLE IN THE 'PROP' FORMAT

1 REFERENCES IN FILE CA (1907 TO DATE)
1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

=> fil capl uspatf toxcenter; s 15
FILE 'CAPLUS' ENTERED AT 12:05:28 ON 08 MAR 2004
USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.
PLEASE SEE "HELP USAGETERMS" FOR DETAILS.
COPYRIGHT (C) 2004 AMERICAN CHEMICAL SOCIETY (ACS)

FILE 'USPATFULL' ENTERED AT 12:05:28 ON 08 MAR 2004
CA INDEXING COPYRIGHT (C) 2004 AMERICAN CHEMICAL SOCIETY (ACS)

FILE 'TOXCENTER' ENTERED AT 12:05:28 ON 08 MAR 2004
COPYRIGHT (C) 2004 ACS

L6 10 L5

*cross Registry answer set into
CAPLUS, USPATFULL, TOXCENTER
to get references*

=> dup rem l6
PROCESSING COMPLETED FOR L6
L7 10 DUP REM L6 (0 DUPLICATES REMOVED)
ANSWERS '1-6' FROM FILE CAPLUS
ANSWER '7' FROM FILE USPATFULL
ANSWERS '8-10' FROM FILE TOXCENTER

=> d ibib ed abs hitstr 1-6; d ibib abs hitstr 7; d iall 8-10; fil hom

L7 ANSWER 1 OF 10 CAPLUS COPYRIGHT 2004 ACS on STN
ACCESSION NUMBER: 2003:818235 CAPLUS
DOCUMENT NUMBER: 139:322283
TITLE: Methods for production and use of mammalian
complementarity determining region mimetibodies for
diagnosis and therapy of human diseases
INVENTOR(S): Heavner, George A.; Knight, David M.; Scallon, Bernard
J.; Ghayeb, John
PATENT ASSIGNEE(S): Centocor, Inc., USA
SOURCE: PCT Int. Appl., 97 pp.
CODEN: PIXXD2
DOCUMENT TYPE: Patent
LANGUAGE: English
FAMILY ACC. NUM. COUNT: 1
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2003084477	A2	20031016	WO 2003-US9139	20030324
W:	AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NI, NO, NZ, OM, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, UZ, VC, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM			
RW:	GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI, SK, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG			

PRIORITY APPLN. INFO.: US 2002-368791P P 20020329

ED Entered STN: 17 Oct 2003

AB This invention pertains to methods for prodn. and use of mammalian complementarity detg. region (CDR) mimetibodies for diagnosis and therapy of human diseases. Genetic engineering, expression, and purifn. of human mimetibodies contg. Ig fragments (CDR, variable, framework and/or const. region) as well as a ligand binding domain are disclosed in this invention. Peptides that mimic the activity of EPO, TPO, growth hormones, G-CSF, GM-CSF, IL-1ra, leptin, CTLA4, TRAIL, TGF-.alpha. and TGF-.beta.

are the focus of this genetic engineering. The aim of the invention is use of the purified recombinant proteins for diagnosis or treatment of anemia, immune or autoimmune disease, cancer, or infectious diseases. At the time of publication, claimed sequence nos. 997 to 1109 were missing, and claimed sequence nos. 984 to 996 were not clearly identified.

IT 148914-10-7

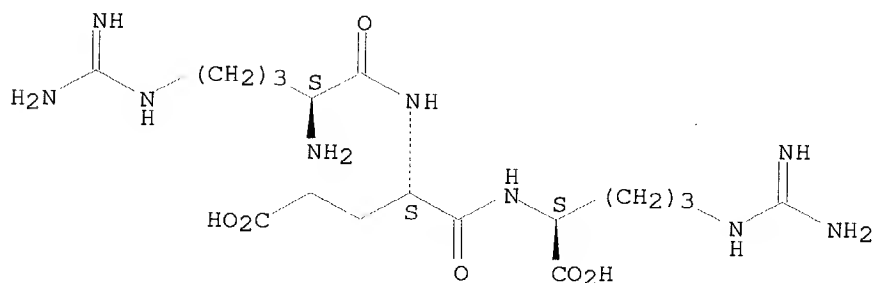
RL: DGN (Diagnostic use); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(macrophage/T cell-inhibiting peptide; methods for prodn. and use of mammalian CDR mimetibodies for diagnosis and therapy of human diseases)

RN 148914-10-7 CAPLUS

CN L-Arginine, L-arginyl-L-.alpha.-glutamyl- (9CI) (CA INDEX NAME)

Absolute stereochemistry.



L7 ANSWER 2 OF 10 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 2002:814182 CAPLUS

DOCUMENT NUMBER: 137:329414

TITLE: Polypeptides for treatment of Alzheimer's disease or use as cognition enhancers

INVENTOR(S): Mileusnic, Radmila; Rose, Steven Peter Russell

PATENT ASSIGNEE(S): The Open University, UK

SOURCE: PCT Int. Appl., 85 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2002083729	A2	20021024	WO 2002-GB1769	20020417
WO 2002083729	A3	20030731		
W:	AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM			
RW:	GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG			
US 2003166529	A1	20030904	US 2001-998491	20011130
EP 1381627	A2	20040121	EP 2002-720228	20020417
R:	AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO, MK, CY, AL, TR			
GB 2391548	A1	20040211	GB 2003-26855	20020417
PRIORITY APPLN. INFO.:			GB 2001-9558	A 20010418

Searched by Barb O'Bryen, STIC 571-272-2518

GB 2001-20084 A 20010817
US 2001-998491 A1 20011130
GB 2002-7387 A 20020328
WO 2002-GB1769 W 20020417

OTHER SOURCE(S): MARPAT 137:329414

ED Entered STN: 25 Oct 2002

AB The invention provides a compd. having formula X1-Arg-Xaa-Arg-X2 in which X1 and X2 are up to 30 amino acid residues and Xaa is an amino acid residue. A preferred compd. is the tripeptide Arg-Glu-Arg which corresponds to amino acid residues 328 to 330 of human amyloid precursor protein. The invention further provides a deriv. of a polypeptide having the formula: X1-Arg-Xaa-Arg-X2 wherein X1 and X2, which may be the same or different, each represents from zero to 30 natural or synthetic amino acid residues or derivs. thereof and Xaa represents a natural or synthetic amino acid residue or deriv. thereof, at least one functional group of at least one said amino acid residue or deriv. thereof being protected by a protective group. The compds. of the invention are believed to be useful in the treatment of Alzheimer's disease and as cognitive enhancers.

IT 148914-10-7

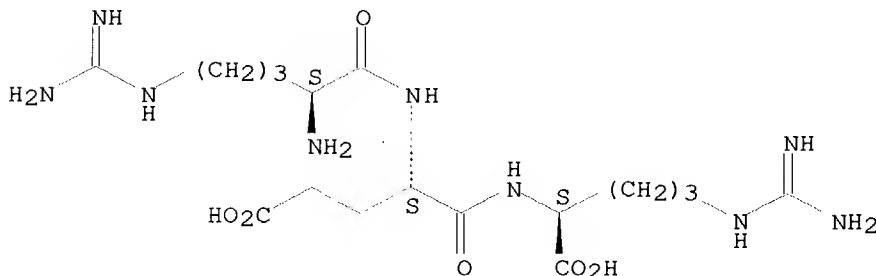
RL: PRP (Properties); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(polypeptides for treatment of Alzheimer's disease or use as cognition enhancers)

RN 148914-10-7 CAPLUS

CN L-Arginine, L-arginyl-L-.alpha.-glutamyl- (9CI) (CA INDEX NAME)

Absolute stereochemistry.



L7 ANSWER 3 OF 10 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER: 1998:175943 CAPLUS

DOCUMENT NUMBER: 128:226237

TITLE: Anti-inflammatory peptides and therapeutic uses thereof

INVENTOR(S): Eisenbach-Schwartz, Michal; Beserman, Pierre; Hirschberg, David L.

PATENT ASSIGNEE(S): Yeda Research and Development Co. Ltd., Israel; Eisenbach-Schwartz, Michal; Beserman, Pierre; Hirschberg, David L.

SOURCE: PCT Int. Appl., 44 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 3

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9809985	A2	19980312	WO 1997-IL295	19970903
WO 9809985	A3	19980507		

Searched by Barb O'Bryen, STIC 571-272-2518

W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
RW: GH, KE, LS, MW, SD, SZ, UG, ZW, AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG

US 6126939 A 20001003 US 1997-864301 19970528
AU 9740301 A1 19980326 AU 1997-40301 19970903
EP 927191 A2 19990707 EP 1997-937794 19970903

R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO

JP 2001500492 T2 20010116 JP 1998-512435 19970903

PRIORITY APPLN. INFO.:

US 1996-25376P P 19960903
US 1996-753141 A 19961120
US 1997-864301 A 19970528
US 1996-31191P P 19961120
WO 1997-IL295 W 19970903

OTHER SOURCE(S): MARPAT 128:226237

ED Entered STN: 25 Mar 1998

AB The invention is directed to peptides of the formulas (i) Xaa-Yaa-Arg (either Xaa is any amino acid residue and Yaa is Glu or Xaa is absent and Yaa is any amino acid residue with the exception of Pro), (ii) Arg-Yaa-Xaa (either Xaa is any amino acid residue and Yaa is Glu or Xaa is absent and Yaa is any amino acid residue with the exception of Asn), (iii) Xaa-Arg-Yaa (Xaa is any amino acid residue and Yaa is Glu), and (i.v.) Yaa-Arg-Xaa (Xaa is any amino acid residue and Yaa is Glu), and to derivs. thereof, which exert an inhibitory effect on macrophage migration and/or macrophage phagocytic activity. In addn., the peptides and derivs. thereof exert an inhibitory effect on the ability of macrophages and T cells to adhere to extracellular matrix and/or fibronectin. The peptides and derivs. thereof exert an inhibitory effect on a humoral and/or cellular immune response. The invention is also directed to methods for use of the peptides and derivs. thereof and compns. contg. them for the inhibition of inflammation, including but not limited to, inflammation at a joint, in the central nervous system generally, at specific lesions in the central nervous system, and other immune privileged sites. Immune privilege factor was purified from brain conditioned medium and shown to have a similar migration pattern to Glu-Arg.

IT 148914-10-7

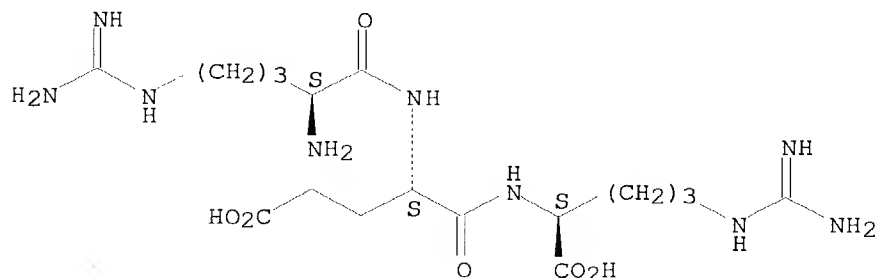
RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(anti-inflammatory peptides and therapeutic uses)

RN 148914-10-7 CAPLUS

CN L-Arginine, L-arginyl-L-.alpha.-glutamyl- (9CI) (CA INDEX NAME)

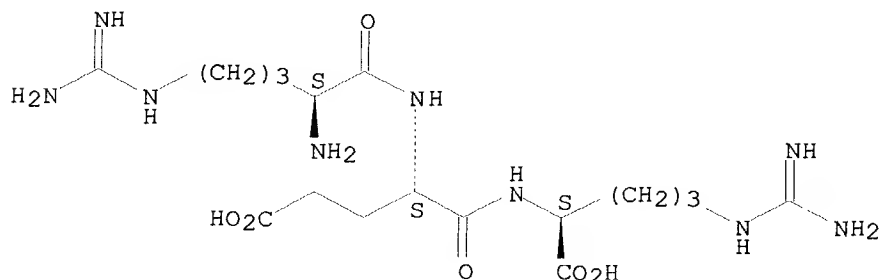
Absolute stereochemistry.



L7 ANSWER 4 OF 10 CAPLUS COPYRIGHT 2004 ACS on STN
ACCESSION NUMBER: 1994:570570 CAPLUS
DOCUMENT NUMBER: 121:170570
TITLE: Substances having the growth-promoting effect of
amyloid precursor protein
INVENTOR(S): Saitoh, Tsunao
PATENT ASSIGNEE(S): University of California, USA
SOURCE: PCT Int. Appl., 115 pp.
CODEN: PIXXD2
DOCUMENT TYPE: Patent
LANGUAGE: English
FAMILY ACC. NUM. COUNT: 1
PATENT INFORMATION:

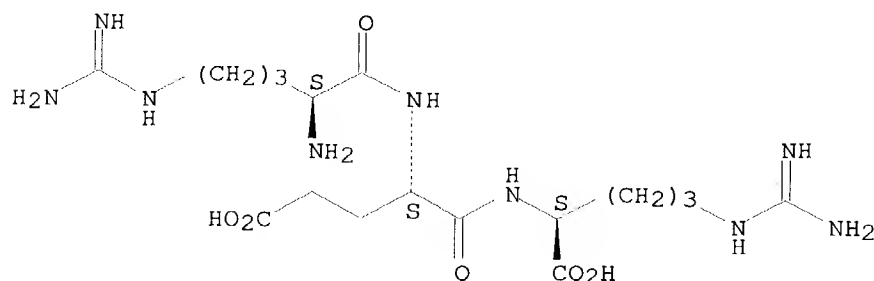
PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9409808	A1	19940511	WO 1992-US9070	19921023
W: AU, CA, JP, KZ, US				
RW: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, SE				
AU 9228951	A1	19940524	AU 1992-28951	19921023
PRIORITY APPLN. INFO.:			WO 1992-US9070	19921023
ED	Entered STN: 15 Oct 1994			
AB	Peptides derived from amyloid precursor protein (APP) that retain at least some neuronal growth promoting effect of APP are described. The peptides include at least five consecutive amino acid residues with side-chain polarities corresponding to the side-chain polarities of the sequence RERMS. Non-peptide compds. with the same activity and methods of prepg. them are described. The peptides and nonpeptides are for use in treatment of neurol. conditions (no data). A series of peptides covering amino acids 296-335 of APP were synthesized and their growth stimulating effects tested on fibroblast cell lines; full-length APP and an analog with a deletion of amino acids 306-335 were prepd. by expression of the cloned gene for use as controls. Only peptides with the RERMS sequence showed growth stimulation and some of the peptides adjacent to the RERMS peptide antagonized its action at high concns. The growth stimulating activity was not due to heparin binding. Studies on the role of APP in neurite outgrowth and sprouting and its interaction with GAP-43 are described. Use of RERMS peptides in the treatment of exptl. spinal ischemia significantly improved the neurol. outcome over the first three days.			
IT	148914-10-7			
	RL: BIOL (Biological study) (peptide of human amyloid precursor protein, growth-promoting properties of)			
RN	148914-10-7 CAPLUS			
CN	L-Arginine, L-arginyl-L-.alpha.-glutamyl- (9CI) (CA INDEX NAME)			

Absolute stereochemistry.



L7 ANSWER 5 OF 10 CAPLUS COPYRIGHT 2004 ACS on STN
ACCESSION NUMBER: 1993:492151 CAPLUS
DOCUMENT NUMBER: 119:92151
TITLE: Amino acid sequence RERMS represents the active domain of amyloid .beta./A4 protein precursor that promotes fibroblast growth
AUTHOR(S): Ninomiya, Haruaki; Roch, Jean Marc; Sundsmo, Mary P.; Otero, Deborah A. C.; Saitoh, Tsunao
CORPORATE SOURCE: Dep. Neurosci., Univ. California, San Diego, La Jolla, CA, 92093, USA
SOURCE: Journal of Cell Biology (1993), 121(4), 879-86
CODEN: JCLBA3; ISSN: 0021-9525
DOCUMENT TYPE: Journal
LANGUAGE: English
ED Entered STN: 04 Sep 1993
AB The growth of A-1 fibroblasts depends on exogenous amyloid .beta./A4 protein precursor (APP), providing a simple bioassay to study the function of APP. To further characterize the growth-promoting activity of the secreted form of APP-695 (sAPP-695) on fibroblasts, the authors applied a battery of synthetic peptides and found that: (1) the sequence of 5 amino acids, RERMS (APP328-332), was uniquely required for the growth-promoting activity of sAPP-695; (2) the activity was sequence specific because the reverse-sequence peptide of the active domain had no activity; and (3) the 4-amino-acid peptide RMSQ (APP330-333), which partially overlaps the C-terminal side of the active sequence RERMS, could antagonize the activity of sAPP-695. Furthermore, a recombinant protein which lacks this active domain (APP20-591 without 306-335) did not promote fibroblast cell growth, suggesting that this domain is the only site of sAPP-695 involved in the growth stimulation. The availability of these biol. active, short peptides and their antagonists should prove to be an essential step for the elucidation of APP involvement in regulation of cellular homeostasis.
IT 148914-10-7
RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study) (fibroblast growth stimulation by, amyloid .beta./A4 protein precursor structure in relation to)
RN 148914-10-7 CAPLUS
CN L-Arginine, L-arginyl-L-.alpha.-glutamyl- (9CI) (CA INDEX NAME)

Absolute stereochemistry.



L7 ANSWER 6 OF 10 CAPLUS COPYRIGHT 2004 ACS on STN
ACCESSION NUMBER: 1987:61216 CAPLUS
DOCUMENT NUMBER: 106:61216
TITLE: Immunoregulatory peptides
INVENTOR(S): Hahn, Gary Scott
PATENT ASSIGNEE(S): Merck Patent G.m.b.H., Fed. Rep. Ger.
SOURCE: PCT Int. Appl., 109 pp.
CODEN: PIXXD2

Searched by Barb O'Bryen, STIC 571-272-2518

DOCUMENT TYPE: Patent
LANGUAGE: English
FAMILY ACC. NUM. COUNT: 1
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE	
WO 8604334	A1	19860731	WO 1986-EP12	19860115	
W: AU, JP					
RW: AT, BE, CH, DE, FR, GB, IT, LU, NL, SE					
AU 8653198	A1	19860813	AU 1986-53198	19860115	
AU 602483	B2	19901018			
EP 215805	A1	19870401	EP 1986-900764	19860115	
R: AT, BE, CH, DE, FR, GB, IT, LI, NL, SE					
JP 62501502	T2	19870618	JP 1986-500766	19860115	
PRIORITY APPLN. INFO.:				US 1985-692711	19850118
				US 1985-803452	19851129
				US 1985-805504	19851129
				WO 1986-EP12	19860115

ED Entered STN: 07 Mar 1987

AB Immunoregulatory peptides AX(BY)nC (X and Y = amino acid residue with pos. charged side chains; A and C = substituents that preserve or augment the immunoregulatory activity of the peptide; B = amino acid residue that preserves or augments the immunoregulatory activity of the peptide; n = 0, 1) are prepd. for use as medicaments for immune system response control. Thus, the bis-trifluoroacetate salt of L-Lys-L-Ser-OH was prepd. by reacting L-serine with N,N'-bis-tert-butyloxycarbonyl-L-lysine N-hydroxysuccinimide ester in THF, deprotection, and reaction with anhyd. trifluoroacetic acid.

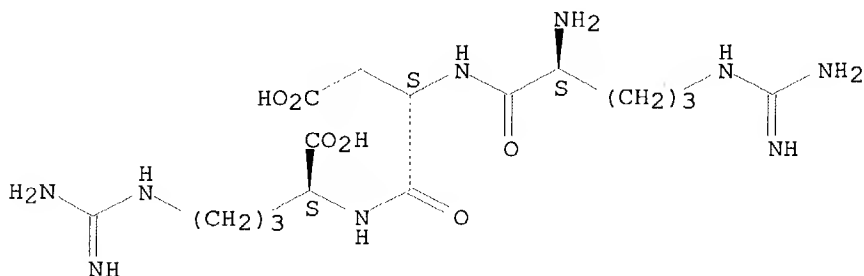
IT 106326-11-8P

RL: SPN (Synthetic preparation); PREP (Preparation)
(prepn. of, or immunoregulator)

RN 106326-11-8 CAPLUS

CN L-Arginine, N2-(N-L-arginyl-L-.alpha.-aspartyl)- (9CI) (CA INDEX NAME)

Absolute stereochemistry.



L7 ANSWER 7 OF 10 USPATFULL on STN

ACCESSION NUMBER: 2003:238371 USPATFULL

TITLE: Polypeptides and their uses

INVENTOR(S): Mileusnic, Radmila, Milton Keynes, UNITED KINGDOM
Russell Rose, Steven Peter, Milton Keynes, UNITED KINGDOM

NUMBER KIND DATE

Searched by Barb O'Bryen, STIC 571-272-2518

PATENT INFORMATION: US 2003166529 A1 20030904
APPLICATION INFO.: US 2001-998491 A1 20011130 (9)

	NUMBER	DATE
PRIORITY INFORMATION:	GB 2001-9558	20010418
	GB 2001-20084	20010817
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	DRINKER BIDDLE & REATH, ONE LOGAN SQUARE, 18TH AND CHERRY STREETS, PHILADELPHIA, PA, 19103-6996	
NUMBER OF CLAIMS:	35	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	7 Drawing Page(s)	
LINE COUNT:	1091	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.
AB The invention provides compounds having formulae comprising

X.sub.1-Ser-Met-Arg-Glu-Arg-X.sub.2

in which X.sub.1 and X.sub.2 are up to 30 amino acid residues and the formula represents a reverse-order sequence corresponding to amino acid residues 332 to 328 of human APP and from zero to 30 successive amino acid residues of human APP extending in each direction therefrom, The invention also provides the pentapeptide Ser-Met-Arg-Glu-Arg, corresponding to residues 332 to 328 of human amyloid precursor protein in reverse order, and the tripeptide Arg-Glu-Arg which corresponds residues 328 to 330 of human amyloid precursor protein, the tripeptide and pentapeptide being provided as pharmaceutical compositions. The invention further provides conjugates of the foregoing compounds which can cross the blood-brain barrier and pharmaceutical compositions containing such conjugates. The compounds and compositions of the invention are believed to be useful in the treatment of Alzheimer's disease and as cognitive enhancers and appropriate methods of medical treatment are disclosed.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

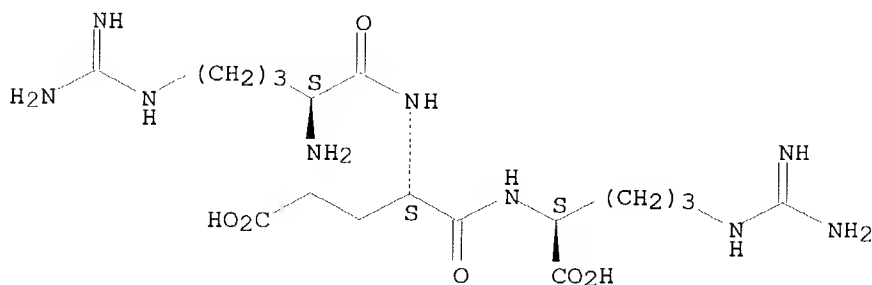
IT 148914-10-7

(polypeptides for treatment of Alzheimer's disease or use as cognition enhancers)

RN 148914-10-7 USPATFULL

CN L-Arginine, L-arginyl-L-.alpha.-glutamyl- (9CI) (CA INDEX NAME)

Absolute stereochemistry.



ACCESSION NUMBER: 2002:119433 TOXCENTER
COPYRIGHT: Copyright 2004 ACS
DOCUMENT NUMBER: CA12819226237F
TITLE: Anti-inflammatory peptides and therapeutic uses thereof
AUTHOR(S): Eisenbach-Schwartz, Michal; Beserman, Pierre; Hirschberg, David L.
CORPORATE SOURCE: ASSIGNEE: Hirschberg, David L.
PATENT INFORMATION: WO 989985 A2 12 Mar 1998
SOURCE: (1998) PCT Int. Appl., 44 pp.
CODEN: PIXXD2.
COUNTRY: ISRAEL
DOCUMENT TYPE: Patent
FILE SEGMENT: CAPLUS
OTHER SOURCE: CAPLUS 1998:175943
LANGUAGE: English
ENTRY DATE: Entered STN: 20020528
Last Updated on STN: 20020605

ABSTRACT:

The invention is directed to peptides of the formulas (i) Xaa-Yaa-Arg (either Xaa is any amino acid residue and Yaa is Glu or Xaa is absent and Yaa is any amino acid residue with the exception of Pro), (ii) Arg-Yaa-Xaa (either Xaa is any amino acid residue and Yaa is Glu or Xaa is absent and Yaa is any amino acid residue with the exception of Asn), (iii) Xaa-Arg-Yaa (Xaa is any amino acid residue and Yaa is Glu), and (i.v.) Yaa-Arg-Xaa (Xaa is any amino acid residue and Yaa is Glu), and to derivs. thereof, which exert an inhibitory effect on macrophage migration and/or macrophage phagocytic activity. In addn., the peptides and derivs. thereof exert an inhibitory effect on the ability of macrophages and T cells to adhere to extracellular matrix and/or fibronectin. The peptides and derivs. thereof exert an inhibitory effect on a humoral and/or cellular immune response. The invention is also directed to methods for use of the peptides and derivs. thereof and compns. contg. them for the inhibition of inflammation, including but not limited to, inflammation at a joint, in the central nervous system generally, at specific lesions in the central nervous system, and other immune privileged sites. Immune privilege factor was purified from brain conditioned medium and shown to have a similar migration pattern to Glu-Arg.

CLASSIFICATION CODE: 1-7

SUPPLEMENTARY TERMS: Miscellaneous Descriptors
immunosuppressant antiinflammatory peptide; immune
privilege factor isolation

REGISTRY NUMBER: 1238-09-1Q (derivs.)
2418-74-8Q (Prolylarginine, derivs.)
2478-01-5Q (derivs.)
2640-07-5Q (derivs.)
7219-59-2Q (derivs.)
13261-11-5Q (Serylarginine, derivs.)
13448-26-5Q (derivs.)
15483-27-9Q (Arginylarginine, derivs.)
16709-12-9Q (derivs.)
18635-55-7Q (Glycylarginine, derivs.)
26607-15-8Q (derivs.)
29586-66-1Q (derivs.)
37682-75-0Q (Valylarginine, derivs.)
55715-01-0Q (derivs.)
60461-10-1Q (derivs.)
68040-95-9Q (derivs.)
70904-56-2Q (derivs.)
77369-21-2Q (derivs.)
88831-09-8Q (derivs.)
126590-89-4Q (derivs.)
186761-64-8Q (derivs.)
204644-00-8Q (derivs.)
204866-55-7 (Immune Privilege Factor)

REGISTRY NUMBER: 1188-24-5; 2047-13-4; 2418-67-9; 2418-69-1; 2483-17-2;
2639-45-4; 2896-20-0; 6418-86-6; 15706-88-4; 15706-89-5;
25615-38-7; 40968-45-4; 40968-46-5; 45243-23-0;
61192-07-2; 62632-70-6; 70921-62-9; 74863-12-0;
82261-72-1; 105425-96-5; 106326-78-7; 115945-15-8;
116685-16-6; 116854-12-7; 125557-81-5; 128500-64-1;
131837-03-1; 132105-44-3; 137427-66-8; 140360-47-0;
140716-02-5; 140716-03-6; 140716-04-7; 140716-05-8;
140716-06-9; 140716-07-0; **148914-10-7**;
155114-05-9; 172684-37-6; 175175-18-5; 175175-24-3;
175175-25-4; 175175-40-3; 175175-41-4; 175175-42-5;
175175-43-6; 175175-44-7; 175175-45-8; 175175-46-9;
175175-47-0; 175175-48-1; 175175-65-2; 175175-91-4;
175176-43-9; 175176-92-8; 175177-31-8; 175177-64-7;
175276-10-5; 182295-52-9; 201463-77-6; 201463-84-5;
204644-01-9; 204644-02-0; 204644-03-1; 204644-04-2;
204644-05-3; 204644-06-4; 204644-07-5; 204644-08-6;
204644-09-7; 204644-10-0; 204644-11-1; 204644-12-2;
204644-13-3; 204644-14-4; 204644-15-5; 204644-16-6;
204644-17-7; 204644-18-8; 204644-19-9; 204644-20-2;
204644-21-3; 204644-22-4; 204644-23-5; 204644-24-6;
204644-25-7; 204644-26-8; 204644-27-9; 204644-28-0;
204644-29-1; 204644-30-4; 204644-31-5; 204644-32-6;
204644-33-7; 204644-34-8

L7 ANSWER 9 OF 10 TOXCENTER COPYRIGHT 2004 ACS on STN
ACCESSION NUMBER: 1994:171025 TOXCENTER
COPYRIGHT: Copyright 2004 ACS
DOCUMENT NUMBER: CA12115170570V
TITLE: Substances having the growth-promoting effect of amyloid
precursor protein
AUTHOR(S): Saitoh, Tsunao
CORPORATE SOURCE: ASSIGNEE: University of California
PATENT INFORMATION: WO 949808 A1 11 May 1994
SOURCE: (1994) PCT Int. Appl., 115 pp.
CODEN: PIXXD2.
COUNTRY: UNITED STATES
DOCUMENT TYPE: Patent
FILE SEGMENT: CAPLUS
OTHER SOURCE: CAPLUS 1994:570570
LANGUAGE: English
ENTRY DATE: Entered STN: 20011116
Last Updated on STN: 20020910

ABSTRACT:

Peptides derived from amyloid precursor protein (APP) that retain at least some neuronal growth promoting effect of APP are described. The peptides include at least five consecutive amino acid residues with side-chain polarities corresponding to the side-chain polarities of the sequence RERMS. Non-peptide compds. with the same activity and methods of prepg. them are described. The peptides and nonpeptides are for use in treatment of neurol. conditions (no data). A series of peptides covering amino acids 296-335 of APP were synthesized and their growth stimulating effects tested on fibroblast cell lines; full-length APP and an analog with a deletion of amino acids 306-335 were prepd. by expression of the cloned gene for use as controls. Only peptides with the RERMS sequence showed growth stimulation and some of the peptides adjacent to the RERMS peptide antagonized its action at high concns. The growth stimulating activity was not due to heparin binding. Studies on the role of APP in neurite outgrowth and sprouting and its interaction with GAP-43 are described. Use of RERMS peptides in the treatment of exptl. spinal ischemia significantly improved the neurol. outcome over the first three days.

CLASSIFICATION CODE: 1-11

SUPPLEMENTARY TERMS: Miscellaneous Descriptors
amyloid precursor growth promoting peptide; neurite

outgrowth amyloid precursor peptide
REGISTRY NUMBER: 148914-08-3Q (peptide and non-peptide analogs)
117-89-5 (Trifluoperazine)
77086-22-7 (MK801)
78990-62-2 (Calpain)
REGISTRY NUMBER: 148914-13-0; 148914-08-3; 157622-71-4; 50-53-3;
148913-99-9; 148914-00-5; 148914-01-6; 148914-02-7;
148914-03-8; 148914-04-9; 148914-05-0; 148914-06-1;
148914-07-2; 148914-09-4; **148914-10-7**;
148914-11-8; 148914-12-9; 149146-19-0

L7 ANSWER 10 OF 10 TOXCENTER COPYRIGHT 2004 ACS on STN
ACCESSION NUMBER: 1987:110406 TOXCENTER
COPYRIGHT: Copyright 2004 ACS
DOCUMENT NUMBER: CA10609061216D
TITLE: Immunoregulatory peptides
AUTHOR(S): Hahn, Gary Scott
CORPORATE SOURCE: ASSIGNEE: Merck Patent G.m.b.H.
PATENT INFORMATION: WO 864334 A1 31 Jul 1986
SOURCE: (1986) PCT Int. Appl., 109 pp.
CODEN: PIXXD2.
COUNTRY: FED. REP. GER.
DOCUMENT TYPE: Patent
FILE SEGMENT: CAPLUS
OTHER SOURCE: CAPLUS 1987:61216
LANGUAGE: English
ENTRY DATE: Entered STN: 20011116
Last Updated on STN: 20021105

ABSTRACT:

Immunoregulatory peptides AX(BY)nC (X and Y = amino acid residue with pos. charged side chains; A and C = substituents that preserve or augment the immunoregulatory activity of the peptide; B = amino acid residue that preserves or augments the immunoregulatory activity of the peptide; n = 0, 1) are prepd. for use as medicaments for immune system response control. Thus, the bis-trifluoroacetate salt of L-Lys-L-Ser-OH was prepd. by reacting L-serine with N,N'-bis-tert-butyloxycarbonyl-L-lysine N-hydroxysuccinimide ester in THF, deprotection, and reaction with anhyd. trifluoroacetic acid.

CLASSIFICATION CODE: 1-6

SUPPLEMENTARY TERMS: Miscellaneous Descriptors
immunoregulation peptide prepn; antitumor peptide prepn
immunoregulation

REGISTRY NUMBER: 6235-35-4 (Lys-Phe)
2130-96-3Q (resin-bound)
2389-45-9Q (resin-bound)
4530-20-5Q (resin-bound)
7764-95-6Q (resin-bound)
13734-34-4Q (resin-bound)
13836-37-8Q (resin-bound)
15761-39-4Q (resin-bound)
23680-31-1Q (resin-bound)
35899-43-5Q (resin-bound)
47173-80-8Q (resin-bound)
55592-81-9Q (benzhydrylamine resin-bound)
78331-03-0Q (benzhydrylamine resin-bound)
106326-22-1Q (benzhydrylamine resin-bound)
106326-23-2Q (resin-bound)
106326-24-3Q (resin-bound)
REGISTRY NUMBER: 56-45-1; 21160-83-8; 30189-36-7; 106326-27-6; 77235-89-3;
106326-26-5; 106326-29-8; 106326-34-5; 106326-36-7;
77236-16-9; 106326-28-7; 106326-31-2; 32388-19-5;
106400-37-7; 6403-11-8; 7369-79-1; 27780-85-4; 92352-82-4;
100385-49-7; 106325-90-0; 106325-94-4; 106326-15-2;
106326-30-1; 106326-32-3; 106326-33-4; 106326-35-6;

106326-37-8; 106326-38-9; 106326-39-0; 106326-40-3;
106326-41-4; 106326-42-5; 106326-43-6; 106326-44-7;
106326-45-8; 106326-46-9; 106326-47-0; 106326-48-1;
106326-49-2; 106326-50-5; 106326-51-6; 106326-52-7;
106326-53-8; 106326-54-9; 106326-55-0; 106326-56-1;
106326-57-2; 106326-58-3; 106326-59-4; 106326-60-7;
106326-61-8; 106326-62-9; 106326-63-0; 106326-64-1;
106326-65-2; 106326-66-3; 106326-67-4; 106326-68-5;
106326-69-6; 106326-70-9; 106326-71-0; 106326-72-1;
106326-73-2; 106326-74-3; 106326-75-4; 106326-76-5;
106326-77-6; 106326-78-7; 106326-79-8; 106326-80-1;
106326-81-2; 106326-82-3; 106326-83-4; 106326-84-5;
106326-85-6; 106326-86-7; 106326-87-8; 106339-43-9;
106339-44-0; 6665-19-6; 20556-18-7; 21438-60-8;
52766-27-5; 55024-09-4; 66138-71-4; 72815-17-9;
74838-86-1; 106325-88-6; 106325-89-7; 106325-91-1;
106325-92-2; 106325-93-3; 106325-95-5; 106325-96-6;
106325-97-7; 106325-98-8; 106325-99-9; 106326-00-5;
106326-01-6; 106326-02-7; 106326-03-8; 106326-04-9;
106326-05-0; 106326-06-1; 106326-07-2; 106326-08-3;
106326-09-4; 106326-10-7; **106326-11-8;**
106326-12-9; 106326-13-0; 106326-14-1; 106326-16-3;
106326-17-4; 106326-18-5; 106326-19-6; 106326-20-9;
106326-21-0; 106339-42-8; 2389-45-9; 2480-93-5; 4530-20-5;
6404-29-1; 7536-58-5; 13139-15-6; 13139-16-7; 13574-13-5;
13734-34-4; 13734-41-3; 13836-37-8; 15260-10-3;
15761-38-3; 15761-39-4; 22838-58-0; 35899-43-5;
55878-47-2; 106326-25-4

FILE 'HOME' ENTERED AT 12:06:14 ON 08 MAR 2004